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    Issued_Patents_NA:*

1: /EMC_Celerra_SIDSJ/ptodata/2/ina/5_COMB.seq:*

2: /EMC_Celerra_SIDSJ/ptodata/2/ina/6A_COMB.seq:*

3: /EMC_Celerra_SIDSJ/ptodata/2/ina/6A_COMB.seq:*

4: /EMC_Celerra_SIDSJ/ptodata/2/ina/6B_COMB.seq:*

5: /EMC_Celerra_SIDSJ/ptodata/2/ina/H_COMB.seq:*

6: /EMC_Celerra_SIDSJ/ptodata/2/ina/H_COMB.seq:*

7: /EMC_Celerra_SIDSJ/ptodata/2/ina/PCTUS_COMB.seq:*

8: /EMC_Celerra_SIDSJ/ptodata/2/ina/PCOMB.seq:*

9: /EMC_Celerra_SIDSJ/ptodata/2/ina/BCCOMB.seq:*

10: /EMC_Celerra_SIDSJ/ptodata/2/ina/BCCOMB.seq:*
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Sequence 100, App
Sequence 98, Appl
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Sequence 95, Appl
Sequence 95, Appl
Sequence 97, Appl
Sequence 97, Appl
Sequence 97, Appl
Sequence 11, Appl
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RESULT 2
US-09-874-601-101
; Sequence 101, Application US/09874601
; Patent No. 6632057
; GENERAL INFORMATION:
; APPLICANT: LEWIN, ALFRED S.

문 S

Matches Best Local Query Match

34;

Similarity

100.0%; Score 34; DE ilarity 100.0%; Pred. No. 3.1 Conservative 0; Mismatches

3.1e-10; DB 3;

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| SULT 1 Sequence 100 Patent No. 6 GENERAL INFO APPLICANT: APPLICANT: TITLS OF IN FILE REFERE CURRENT APPL PRIOR APPLI PRIOR PILIN OUMBER OF S SOFTWARE: PRIOR TYPE: RNA ORGANISM: TYPE: RNA ORGANISM: PEATURE: NAME/KEY: INGCATION: OTHER INFO  |            |   |   |
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| RESULT 1  US-09-874-601-100  ¡ Sequence 100, Application ¡ Patent No. 6632057 ¡ GENERAL INFORMATION: ¡ APPLICANT: LEWIN, ALFRED APPLICANT: SHAW, LYNN C APPLICANT: GRANT, MARIA TITLE OF INVENTION: ADEN ¡ TITLE OF INVENTION: THE FILE REFERENCE: 4300.014 ¡ CURRENT APPLICATION NUMBER CURRENT FILING DATE: 20 PRIOR APPLICATION NUMBER PRIOR FILING DATE: 1997- PRIOR PILING DAT   |            |   |   |
| .00 Appli Appli Appli SHAWIN, GRANTION GRANT, FINTION DATE: ATION DATE: ATION DATE: ATION CENTION CONTROL ATION CO   |            | 444444  | 44444444444   |
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| OP-874-601-100 equence 100, Application US atent No. 6632057 satent No. 6632057 satent No. 6632057 sapplicant: LEWIN, ALFRED S. APPLICANT: LEWIN, ALFRED S. APPLICANT: GRANT, MARIA B. TITLE OF INVENTION: ADENO-A FILE REFERENCE: 4300.014100 CURRENT APPLICATION NUMBER: CURRENT FILING DATE: 2001- PRIOR FILING DATE: 1998-05- PRIOR FILING DATE: 1997-05- PRIOR APPLICATION NUMBER: 6 PRIOR PILING DATE: 1997-05- PRIOR APPLICATION NUMBER: 6 PRIOR FILING DATE: 1997-05- PRIOR APPLICATION NUMBER: 6 PRIOR PILING DATE: 1997-05- PRIOR APPLICATION NUMBER: 6 PRIOR APPLICATION NUMBER: 6 PRIOR FILING DATE: 1997-05- PRIOR APPLICATION NUMBER: 6 PRIOR FILING DATE: 1997-05- PRIOR FILING DATE: 1997-05- PRIOR PRIOR FILING DATE: 1997-05- PRIOR FILI   |            | 38<br>38<br>491<br>498<br>570<br>601  | 1605<br>1605<br>1605<br>1991<br>1991<br>1991<br>12624<br>2624<br>2624<br>2671<br>125192<br>125192<br>276687<br>125192<br>44103765<br>44103765   |
| SULT 1  Sequence 100, Application US/09874601  Sequence 100, Application US/09874601  Sequence 100, Application US/09874601  Patent No. 6632057  GENERAL INFORMATION: APPLICANT: LEWIN, ALFRED S. APPLICANT: SHAW, LYNN C. APPLICANT: GRANT, MARIA B.  TITLE OF INVENTION: ADENO-ASSOCIATED VIRUS-DELIVERED RIBOZYME  TITLE OF INVENTION: THE TREATMENT OF RETINAL DISEASES  FILE REFERENCE: 4300.014100  CURRENT SPLICATION NUMBER: US/09/874,601  CURRENT FILING DATE: 2001-05-01  PRIOR APPLICATION NUMBER: 60/046,147  PRIOR FILING DATE: 1997-05-09  PRIOR APPLICATION NUMBER: 60/044,492  PRIOR FILING DATE: 1997-04-21  NUMBER OF SEQ ID NOS: 182  SOFTWARE: Patentin version 3.0  SEQ ID NO 100  LENGTH: 34  TYPE: RNA ORGANISM: Artificial Sequence  PEATURE: NAME/KEY: misc feature LOCATION: ()7)  OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE  -09-874-601-100  |            | <b></b>   |   |
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| 74601<br>ATED<br>ENT 0<br>5,667<br>1,492   | ž          | US-09-<br>US-09-<br>US-09-<br>US-09-<br>US-09-  | US-10-<br>US-10-<br>US-10-<br>US-09-<br>US-08-<br>US-08-<br>US-08-<br>US-09-<br>US-09-<br>US-09-<br>US-09-<br>US-09-<br>US-09-<br>US-09-  |
| )1<br>SD V<br>OF V<br>974,   | ALIGNMENTS | US-10-138-674B-1362<br>US-10-138-674B-14209<br>US-09-640-211A-529<br>US-09-902-540-8002<br>US-09-328-352-2408<br>US-09-349-016-121350<br>US-09-533-559-3649 | US-10-291-446-1 US-10-241-486-3 US-10-244-851-3 US-09-270-767-11487 US-09-232-144-3 US-08-232-144-5 US-08-232-144-5 US-08-99-9016-14120 US-08-99-9016-1840A-2 US-09-371-7728-1158   |
| 71 RE 80 3   | ME         | 138-674B-13866<br>138-674B-13866<br>138-674B-1409<br>640-211A-529<br>902-540-8002<br>328-352-2408<br>949-016-121350<br>533-559-3649                         | 291-446-1<br>410-648-3<br>274-851-3<br>270-767-114<br>232-144-5<br>708-392-10<br>232-144-5<br>232-144-5<br>232-144-5<br>1949-016-1-11<br>949-016-1-11<br>949-016-1-13<br>949-016-1-13<br>34-847-806   |
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| SIT  |            | 11366,<br>113866,<br>14209, Ap<br>529, Ap<br>8002, A<br>2408, A<br>121350,<br>3649, A   | 1, Appl<br>3, Appl<br>3, Appl<br>11487,<br>10, Appl<br>10, Appl<br>11, Appl<br>11, Appl<br>11, Appl<br>11, Appl<br>11, Appl<br>11, Appl<br>11, Appl<br>11, Appl<br>11, Appl   |
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CURRENT APPLICATION NUMBER: US/09/874,601
CURRENT FILING DATE: 2001-05-01
PRIOR APPLICATION NUMBER: 09/063,667
PRIOR FILING DATE: 1998-04-21
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR FILING DATE: 1997-05-09
PRIOR FILING DATE: 1997-04-21
PRIOR APPLICATION NUMBER: 60/044,492
PRIOR FILING DATE: 1997-04-21
NUMBER OF SEQ ID NOS: 182
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Patent No. 6
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APPLICANT: SHAW, LYNN C.
APPLICANT: GRANT, MARIA B.
APPLICANT: GRANT, MARIA B.
TITLE OF INVENTION: ADENO-ASSOCIATED VIRUS-DELIVERED RIBOZYME COMPOSITIONS AND METHOD TITLE OF INVENTION: THE TREATMENT OF RETINAL DISEASES
FILE REFERENCE: 4300.014100
CURRENT APPLICATION NUMBER: US/09/874,601
CURRENT FILING DATE: 2001-05-01
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PRIOR FILING DATE: 1997-05-09
PRIOR APPLICATION NUMBER: 60/044,492
PRIOR FILING DATE: 1997-04-21
NUMBER OF SEQ ID NOS: 182
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NAME/KEY: misc feature
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OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE
09-874-601-101
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ORCANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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GRANT, MARIA B.
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US-09-874-601-93
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SEQ ID NO 92
LENGTH: 34
TYPE: RNA
ORGANISM: Artificial Sequence
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Patent No. 6632057
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                     APPLICANT: LEWIN, ALFRED S.
APPLICANT: SHAW, LYNN C.
APPLICANT: GRANT, MARIA B.

APPLICANT: GRANT, MARIA B.

TITLE OF INVENTION: ADENO-ASSOCIATED VIRUS-DELIVERED RIBOZYME COMPOSITIONS AND METHODS
FILE REFERENCE: 4300.014100

CURRENT APPLICATION NUMBER: US/09/874,601

CURRENT FILING DATE: 2001-05-01

PRIOR APPLICATION NUMBER: 60/046,147

PRIOR APPLICATION NUMBER: 60/046,147

PRIOR APPLICATION NUMBER: 60/046,147

PRIOR APPLICATION NUMBER: 60/046,147

PRIOR FILING DATE: 1997-05-09

PRIOR FILING DATE: 1997-05-09
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TITLE OF INVENTION: THE TREATMENT OF RETINAL DISEASES
FILE REFERENCE: 4300.014100
CURRENT APPLICATION UNMER: US/09/874,601
CURRENT FILING DATE: 2001-05-01
PRIOR APPLICATION NUMBER: 09/063,667
PRIOR PILING DATE: 1998-04-21
PRIOR APPLICATION NUMBER: 60/044,147
PRIOR PILING DATE: 1997-05-09
PRIOR APPLICATION NUMBER: 60/044,492
PRIOR APPLICATION NUMBER: 60/044,492
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NUMBER OF SEQ ID NOS: 182
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             NAME/KEY: misc_feature
LOCATION: ()..()
OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE
                                                                                                    TYPE: RNA ORGANISM: Artificial Sequence FEATURE:
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NAME/KEY: misc_feature
LOCATION: ()..()
OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE
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100.0%; Pr
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GENERAL INFORMATION:
APPLICANT: LEWIM, ALFRED S.
APPLICANT: SHAW, LYNN C.
APPLICANT: SHAW, LYNN C.
APPLICANT: GRANT, MARIA B.
TITLE OF INVENTION: ADENO-ASSOCIATED VIRUS-DELIVERED RIBOZYME COMPOSITIONS AND METHOD
TITLE OF INVENTION: THE TREATMENT OF RETINAL DISEASES
FILE REFERENCE: 4300.014100
CURRENT APPLICATION NUMBER: US/09/874,601
CURRENT FILING DATE: 09/063.667
PRIOR APPLICATION NUMBER: 09/063,667
PRIOR FILING DATE: 1998-04-21
PRIOR FILING DATE: 1998-04-21
PRIOR FILING DATE: 1997-04-21
NUMBER: OF SEQ ID NOS: 182
SOPTWARE: Patentin version 3.0
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CURRENT FILING DATE: 2001-05-01
PRIOR APPLICATION NUMBER: 09/063,667
PRIOR FILING DATE: 1998-04-21
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR FILING DATE: 1997-05-09
PRIOR APPLICATION NUMBER: 60/044,492
PRIOR APPLICATION NUMBER: 60/044,492
PRIOR PILING DATE: 1997-04-21
NUMBER OF SEQ ID NOS: 182
SOPTWARE: PATENTIN VETSION 3.0
SEQ ID NO 99
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Patent No. 6632057
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APPLICANT: LEWIN, ALFRED S.
APPLICANT: SHAW, LYNN C.
APPLICANT: GRANT, MARIA B.
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TITLE OF INVENTION: THE TREATMENT OF RETINAL DISEASES
FILE REFERENCE: 4300.014100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: RNA
ORGANISM: Artificial Sequence
FEATURE:
NAME/KEY: misc_feature
LOCATION: ()..()
OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE
ORGANISM: Artificial Sequence
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Patent No. 6632057
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Best Local Similarity
                                                                                                                                                                                                                                                                                                 Patent No.
APPLICANT: LEWIN, ALFRED S.
APPLICANT: STANW, LYNN C.
APPLICANT: GRAW, LYNN C.
APPLICATION: ADENO-ASSOCIATED VIRUS-DELIVERED RIBOZYME COMPOSITIONS AND METHODS:
TITLE OF INVENTION: THE TREATMENT OF RETINAL DISEASES
FILE REPERENCE: 4300.014100
CURRENT APPLICATION NUMBER: US/09/874,601
CURRENT FILING DATE: 2001-05-01
PRIOR APPLICATION NUMBER: 09/063,667
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR PILING DATE: 1998-04-21
FRIOR FILING DATE: 1997-05-09
PRIOR FILING DATE: 1997-05-09
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TITLE OF INVENTION: THE TREATMENT OF RETINAL DISEASES
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PRIOR APPLICATION NUMBER: 60/044,492
PRIOR APPLICATION NUMBER: 60/044,492
PRIOR PILING DATE: 1997-04-21
NUMBER: 0F SED ID NOS: 182
COUNTAINE OF SED ID NOS: 182
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APPLICANT: SHAW, LYNN C.
APPLICANT: GRANT, MARIA B.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: RNA
ORGANISM: Artificial Sequence
FEATURE:
NAME/KEY: misc feature
LOCATION: ()...()
OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE
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LOCATION: ()..()
OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE
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100.0%; Pred. No. 4.5e-05;
rative 0; Mismatches 0;
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APPLICATION NUMBER: 60/044,492

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APPLICANT: LEWIN, ALFRED S.

APPLICANT: SHAW, LYNN C.

APPLICANTON: ADENO-ASSOCIATED VIRUS-DELIVERED RIBOZYME COMPOSITIONS AND METHOD TITLE OF INVENTION: ADENO-ASSOCIATED VIRUS-DELIVERED RIBOZYME COMPOSITIONS AND METHOD TITLE OF INVENTION: ADENO-ASSOCIATED VIRUS-BRING OF RETINAL DISEASES

FILE REFERENCE: 4300.014100

CURRENT FILING DATE: 2001-05-01

PRIOR APPLICATION NUMBER: 09/063,667

PRIOR FILING DATE: 1998-04-21

PRIOR FILING DATE: 1997-04-21

PRIOR APPLICATION NUMBER: 60/044,492

PRIOR APPLICATION NUMBER: 60/044,492

PRIOR APPLICATION NUMBER: 60/044,492

PRIOR FILING DATE: 1997-04-21

NUMBER OF SEQ ID NOS: 182

SOFTWARE: Patentin version 3.0

SEQ ID NO 96

LENGTH: 34
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, NAME/KEY: misc feature
; LOCATION: ()..()
; OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE
US-09-874-601-95
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                                                                                                                                   Sequence 97, Application US/09874601 Patent No. 6632057 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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GENERAL INFORMATION:
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APPLICANT: LEWIN, ALFRED S.
APPLICANT: SHAW, LYNN C.
APPLICANT: GRANT, MARIA B.
TITLE OF INVENTION: ADENO-ASSOCIATED VIRUS-DELIVERED RIBOZYME COMPOSITIONS AND METHOD TITLE OF INVENTION: THE TREATMENT OF RETINAL DISEASES
FILE REFERENCE: 4300.014100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 1997-04-21
NUMBER OF SEQ ID NOS: 182
SOFTWARE: PatentIn version 3.0
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LOCATION: ()...()
OTHER INFORMATION: SYNTHETIC OLIGONUCLECTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
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f; Pred. No. 4.5
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LENGTH: 34
TYPE: RNA
CORGANISM: Artificial Sequence
FEATURE:
FEATURE:
LOCATION: ()...()
COTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE
US-09-874-601-94
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US-09-874-601-94
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CURRENT FILING DATE: 2001-05-01
PRIOR APPLICATION NUMBER: 09/063,667
PRIOR FILING DATE: 1998-04-21
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR FILING DATE: 1997-05-09
PRIOR APPLICATION NUMBER: 60/044,492
PRIOR APPLICATION NUMBER: 60/044,492
PRIOR FILING DATE: 1997-04-21
NUMBER OF SEO ID NOS: 182
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn version 3.0 SEQ ID NO 94
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GENERAL INFORMATION:
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LENGTH: 34
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Best Local 9
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TITLE OF INVENTION: THE TREATMENT OF RETINAL DISEASES
FILE REFERENCE: 4300.014100
CURRENT APPLICATION NUMBER: US/09/874,601
CURRENT FILING DATE: 2001-05-01
PRIOR APPLICATION NUMBER: 09/063,667
PRIOR PRIOR DATE: 1998-04-21
PRIOR PRIOR DATE: 1998-04-21
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR FILING DATE: 1997-05-09
PRIOR FILING DATE: 1997-04-21
RIOR FILING DATE: 1997-04-21
RIOR FILING DATE: 1997-04-21
RIOR FILING DATE: 1997-04-21
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APPLICANT: SHAW, LYNN C.
APPLICANT: GRANT, MARIA B.
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OTHER INFORMATION: SYNTHETIC OLIGONUCLECTIDE
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                                     12 GAGCCGUUCGCGGCGAAAC 30
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0; Mismatches
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RESULT 13 US-09-489-039A-3537/c

Sequence 3537, Application US/09489039A Patent No. 6610836

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CURRENT APPLICATION NUMBER: US/09/489,039A; CURRENT FILING DATE: 2000-01-27; PRIOR APPLICATION NUMBER: US 60/117,747; PRIOR FILING DATE: 199-01-29; NUMBER OF SEQ ID NOS: 14342; SEQ ID NO 3537; ELNGTH: 33; TYPE: DNA ORGANISM: Klebsiella pneumoniae
US-09-489-039A-3537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TYPE: DNA
; ORGANISM: Human
US-09-949-016-81813
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR PAPPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR PILING DATE: 2000-09-08
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SOPTWARE: PastSEQ for
SEQ ID NO 81813
LENGTH: 601
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Patent No. 6087154
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Best Local Similarity
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GENERAL INFORMATION:
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APPLICANT: Gary Br
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                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
TILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE:
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
TITLE OF INVENTION: PARUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
                                                      APPLICANT: Baez, Melvyn
APPLICANT: Cates, Carolyn A.
APPLICANT: Cates, Carolyn A.
TITLE OF INVENTION: RHESUS NEUROPEPTIDE Y1 RECEPTOR
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
ADDRESSEE: Eli Lilly and Company
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
              ADDRESSEE: Eli Lilly and Company STREET: Lill Corporate Center CITY: Indianapolis
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Pred. No. 26;
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Pred. No. 26;
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Search completed: August 28, 2006, 10:29:28 Job time : 154 secs

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; LOCATION:
US-09-045-186-1
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                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Gaylo, Paul J.
REGISTRATION NUMBER: 36,808
                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 1152 base pair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 46285
COMPUTER READABLE FORM:
TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: (317) 276-0756
                                                                                                                                                                                             MOLECULE TYPE:
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                                                                               Local
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538 ACTGATGAGCCGTTC 552
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11; Conserv
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seq length: 2000000000
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

| 1<br>2<br>2<br>3<br>3<br>4<br>4<br>4<br>4<br>4<br>7<br>7<br>7<br>10<br>11<br>11<br>11<br>11<br>11<br>11<br>11<br>11<br>11<br>11<br>11<br>11  | Result         |
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| ABZ72865 ABZ72866 ABZ72863 ABZ72857 ABZ72858 ABZ72858 ABZ72864 ABZ72873 ABZ72864 ABZ72865 ABZ72866 ABZ72866 ABZ72866 ABZ72866 ABZ72866 ABZ72860 ABZ72861 AAC88559 AAC88555 AAC88555 AAC88555   | SUMMARIES      |
| Abz72865 IGF1 Rz1 Abz72866 IGF1 Rz2 Abz72867 Alpha 3 R Abz72857 Alpha 3 R Abz72858 Alpha 3 R Abz72858 Alpha 3 R Abz72858 Alpha 3 R Abz72872 VEGFR2 Rz Abz72857 VEGFR2 Rz Abz72855 A2B Rz1 e Abz72855 A2B Rz1 e Abz72856 A2B Rz2 e Abz72860 Alpha 1 R Abz72860 VEGFR1 Rz Abz72860 VEGFR1 Rz Abz72860 Alpha 5 R Abz72861 Alpha 1 R Aac88554 Oligonucl Aac88555 Anti-SOD- Aac88559 Anti-SOD- Aac88559 Anti-SOD- | Description    |

| a                  | _                  |                    |                    |                    | _                 | _                 | _                 | a                  | _                  | _                  | _                 | _                  | _                  | _                  | _                  |                    |                    |                    |                    |                    | _                  | _                  |                    |                    |                    |                    |
|--------------------|--------------------|--------------------|--------------------|--------------------|-------------------|-------------------|-------------------|--------------------|--------------------|--------------------|-------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|
| 45                 | 44                 | 43                 | 42                 | 41                 | 40                | : 39              |                   |                    |                    | c<br>35            |                   | C 33               |                    |                    | : 30               | 29                 | 28                 | 27                 | 26                 | 25                 |                    | c 23               | 22                 | 21                 | 20                 | 19                 |
|                    |                    |                    |                    |                    |                   |                   |                   |                    |                    |                    |                   |                    |                    |                    |                    |                    |                    |                    |                    |                    |                    |                    |                    |                    |                    |                    |
| 15                 | 16                 | 16                 | 16                 | 16                 | 16                | 16                | 16                | 16                 | 16                 | 16                 | 16                | 16                 | 16                 | 16                 | 16                 | 19                 | 24                 | 24                 | 24                 | 24                 | 24                 | 24                 | 24                 | 24                 | 24                 | 24                 |
| 44.                | 47.1               | 47.                | 47.1               | 47.                | 47.1              | 47.1              | 47.1              | 47.1               | 47.1               | 47.1               | 47.1              | 47.1               | 47.1               | 47.1               | 47.1               | 55.9               | 70.6               | 70.6               | 70.6               | 70.6               | 70.6               | 70.6               | 70.6               | 70.6               | 70.6               | 70.                |
| 1                  | _                  | _                  | _                  | -                  |                   | _                 | -                 | щ                  | ш                  | μ                  | щ                 | щ                  | -                  | ۳,                 | _                  | 9                  | σ                  | 0                  | σ                  | 0                  | σ                  | σ                  | σ                  | 0                  | 0                  | σ                  |
| 100                | 6919               | 2496               | 1549               | 1452               | 1143              | 1143              | 1143              | 1143               | 1143               | 1143               | 1140              | 564                | 564                | 564                | 456                | 34                 | 44                 | 44                 | 44                 | 36                 | 36                 | 36                 | 35                 | 35                 | 34                 | 34                 |
| œ                  | 4.                 | 10                 | 9                  | æ                  | 15                | 15                | 15                | 15                 | 15                 | 15                 | 15                | 15                 | 15                 | 15                 | IJ                 | 10                 | w                  | w                  | ω                  | w                  | w                  | w                  | w                  | ω                  | 10                 | ω                  |
| ACD80108           | ABL05593           | ADC51485           | ADA48529           | ADA70476           | AEF16579          | AEF16577          | AEF16578          | AEF16518           | AEF16522           | AEF16520           | AEF16575          | AEF16551           | AEF16552           | AEF16550           | ACF82077           | ABZ72859           | AAC88592           | AAC88596           | AAC88598           | AAC88593           | AAC88597           | AAC88599           | AAC88535           | AAC88537           | ABZ72871           | AAC88561           |
| Acd80108 B. coli K | Abl05593 Drosophil | Adc51485 Bacterial | Ada48529 Rice gene | Ada70476 Rice gene | Aef16579 Alkaline | Aef16577 Alkaline | Aef16578 Alkaline | Aef16518 Nocardiop | Aef16522 Nocardiop | Aef16520 Nocardiop | Aef16575 Alkaline | Aef16551 Nocardiop | Aef16552 Nocardiop | Aef16550 Nocardiop | Acf82077 Human SIR | Abz72859 Alpha 5 R | Aac88592 Oligonucl | Aac88596 Oligonucl | Aac88598 Oligonucl | Aac88593 Oligonucl | Aac88597 Oligonucl | Aac88599 Oligonucl | Aac88535 Anti-gamm | Aac88537 Anti-gamm | Abz72871 VEGFR1 Rz | Aac88561 Anti-SOD- |

## ALIGNMENTS

RESULT 1
ABZ72865
ID 7ABZ7
XX ABZ7
XX ABZ7
XX ABZ7
XX ABZ7
XX Haix
XX Haix
XX Opht
XX Haix
XX Opht
XX Opht
XX OPH
XX OT-N
XX O1-N
XX O Hairpin ribozyme; hammerhead ribozyme; ribozyme; retinal disease; target; ophthalmological; gene therapy; eye; retinal dysfunction; AAV; diabetic retinopathy; mecular degeneration; autosomal dominant retinitis; blood-retinal barrier dysfunction; adeno-associated virus; blindness; ss. 07-NOV-2002. Synthetic. WO200288320-A2. IGF1 Rz1 exemplary ribozyme construct SEQ ID NO:100 09-APR-2003 ABZ72865; ABZ72865 standard; RNA; 34 (first entry) BP.

WPI; 2003-111880/10. 01-MAY-2001; 2001US-00847601. Lewin AS, 01-MAY-2002; 2002WO-US013679. (UYFL ) UNIV FLORIDA. Shaw LC, Grant MB;

Claim 1; Page 81; 115pp; English.

degeneration.

A recombinant adeno-associated virus-vectored ribozyme composition, useful for treating a disease or dysfunction of the mammalian eye e retinal disease, e.g. diabetic retinopathy or age-related macular

e.g.

The present invention describes a recombinant adeno-associated virus (AAV) vectored ribozyme composition (I). (I) comprises: (a) at least first ribozyme that specifically cleaves an mRNA encoding a protein, g

Claim 1; Page 81; 115pp; English.

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                                               A recombinant adeno-associated virus-vectored ribozyme composition, useful for treating a disease or dysfunction of the mammalian eye e retinal disease, e.g. diabetic retinopathy or age-related macular
                                                                                                                                                                                                                                                                                                                                                                                                                                                    ophthalmological; gene therapy; eye; retinal dysfunction; AAV; diabetic retinopathy; macular degeneration; autosomal dominant retinitis; blood-retinal barrier dysfunction; adeno-associated virus; blindness; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hairpin ribozyme;
ophthalmological;
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                                                                                                                                                                       Shaw LC,
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Pred. No. 6.9
0; Mismatches
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RESULT 3
ABZ772863
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CC first ribozyme that specifically cleaves an mRNA encoding a protein, concern the properties, or peptide selected from the group of rod opsin, iNOS, cc RDS/peripherin, VEGFR1, VEGFR2, adenosine A-2B receptor, IGF-1, integrin alpha 1, integrin alpha 3, integrin alpha 5, or integrin alpha 1, concern the comprising a polymucleotide encoding the ribozyme, where the comprising a polymucleotide encoding the ribozyme, where the comprising the ribozyme or the polymucleotide in a selected cc comprising the ribozyme or the polymucleotide; or (e) a host cell comprising the ribozyme or the polymucleotide; or (e) a host cell comprising the ribozyme or the polymucleotide; or (e) a host cell comprising the ribozyme or the polymucleotide. Also described is a method confirmed to mammalian eye, comprising providing to the eye the composition described above, and for a time effective to specifically cell cave the mRNA in the cell. (I) has ophthalmological activity, and can be used in gene therapy. (I) can be used for treating a disease or retinal cell disease mentioned above, including and dominant cell treating the disease mentioned above, including and companient for treating the disease mentioned above, including and companient of a pathological condition, e.g. atrophic or pigmented lesions of the eye companient of providing and condition, e.g. atrophic or pigmented lesions of the eye condition of the mammalian condition, or a reduction in central or peripheral vision, or a reduction in central or peripheral vision, or a reduction in central or peripheral vision, or a reduction in central condition of the providing and central central condition of the providing and central central condition of the providing and central vision of the eye.
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Best Local
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                                    WPI; 2003-111880/10
                                                                                                         Lewin AS,
                                                                                                                                                                                                                                            01-MAY-2001; 2001US-00847601.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    exemplification of the present invention
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V) vectored ribozyme composition (I). (I) comprises: (a) at least a
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                                                                                                     Shaw LC,
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                                                                                                     Grant MB;
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Pred. No.
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The present invention describes a recombinant adeno-associated virus CC (AAV) vectored ribozyme composition (I). (I) comprises: (a) at least a CC first ribozyme that specifically cleaves an mRNA encoding a protein, CC polypeptide, or peptide selected from the group of rod opsin, INOS, CC RDS/peripherin, VEGFR1, VEGFR2, adenosine A-2B receptor, IGF-1, integrin alpha 1, integrin alpha 3, integrin alpha 5, or integrin alpha V; (b) a CC vector comprising a polynucleotide encoding the ribozyme where the CC polynucleotide operably positioned downstream of at least a first CC promoter that directs expression of the polynucleotide in a selected CC mammalian cell transformed with the vector; (c) a viral particle CC comprising the ribozyme or the polynucleotide; (d) an AAV vector CC comprising the ribozyme or the polynucleotide; or (e) a host cell CC comprising the ribozyme or the polynucleotide; or (e) a host cell CC comprising the amount of mRNA encoding a selected polypeptide in a CC composition described above, and for a time effective to specifically CC cleave the mRNA in the cell. (I) has ophthalmological activity, and can be used in gene therapy. (I) can be used for treating a disease or CC dysfunction, (diabetic) retinopathy, or (age-related) macular contraction the mammalian eye, such as a retinal disease or retinal CC reating the disease mentioned above, including autosomal dominant CC creating the disease mentioned above, including autosomal dominant contraction, decreasing the severity, or amelioration the symptoms of a contraction of the mammalian eye, such as a retinal disease or retinal CC contraction and accurating the symptoms of a contraction of the mammalian eye, such as a retinal disease or retinal CC contraction of the mammalian eye, such as a retinal disease or retinal contraction of the mammalian eye, such as a retinal disease or retinal contraction of the mammalian eye, such as a retinal disease or retinal contraction. (I) can also be useful
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Best Local S
Matches 27
                                                                                                                                                                                                                                                                                                                                                                                      Hairpin ribozyme; hammerhead ribozyme; ribozyme; retinal disease; target; ophthalmological; gene therapy; eye; retinal dysfunction; AAV; diabetic retinopathy; macular degeneration; autosomal dominant retinitis; blood-retinal barrier dysfunction; adeno-associated virus; blindness; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for treating, decreasing the severity, or ameliorating the symptoms of a pathological condition, e.g. atrophic or pigmented lesions of the eye, blindness, a reduction in central or peripheral vision, or a reduction in total vision. ABZ72763 to ABZ72953 represent sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A recombinant adeno-associated virus-vectored ribozyme composition, useful for treating a disease or dysfunction of the mammalian eye e.g. retinal disease, e.g. diabetic retinopathy or age-related macular
                                                             01-MAY-2001; 2001US-00847601
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AACUGAUGAGCCGUUCGCGGCGAAACG
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Pred. No.
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4.7e-05;
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(UYFL ) UNIV FLORIDA

07-NOV-2002

WO200288320-A2

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The present invention describes a recombinant adeno-associated virus (AAV) vectored ribozyme composition (I). (I) comprises: (a) at least a C first ribozyme that specifically cleaves an mRNA encoding a protein, copolypeptide, or peptide selected from the group of rod opsin, INOS. CC RDS/peripherin, VEGFR1, VEGFR2, adenosine A-2B receptor, IGF-1, integrin alpha 1, integrin alpha 3, integrin alpha 5, or integrin alpha V; (b) a vector comprising a polymucleotide encoding the ribozyme V; (b) a promoter that directs expression of the polymucleotide in a selected comprising the ribozyme or the polymucleotide; (c) a viral particle comprising the ribozyme or the polymucleotide; (d) an AAV vector comprising the ribozyme or the polymucleotide; or (e) a host cell comprising the ribozyme or the polymucleotide; or (e) a host cell comprising the ribozyme or the polymucleotide; or (e) a host cell composition described above, and for a time effective to specifically cleave the mRNA in the cell. (I) has ophthalmological activity, and can be used in gene therapy. (I) can be used for treating a disease or dysfunction, (diabetic) retinopathy, or (age-related) macular cell dysfunction, (diabetic) retinopathy, or (age-related) macular cell for treating, decreasing the severity, or ameliorating the symptoms of a pathological condition, e.g. atrophic or pigmented lesions of the eye, comprision, or a reduction in central or peripheral vision, or a reduction in central or peripheral vision.
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                                                                                                                                 Hairpin ribozyme; hammerhead ribozyme; ribozyme; retinal disease; target; ophthalmological; gene therapy; eye; retinal dysfunction; AAV; diabetic retinopathy; macular degeneration; autosomal dominant retinitis; black-retinal harrier dysfunction; adeno-associated virus; blindness; 88.
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                                                                                            Synthetic
                                                                                                                                                                                                                                                                             09-APR-2003
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                                                                                                                                                                                                                                 exemplary ribozyme
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                Hairpin ribozyme; hammerhead ribozyme; ribozyme; retinal disease; target; ophthalmological; gene therapy; eye; retinal dysfunction; AAV; diabetic retinopathy; macular degeneration; autosomal dominant retinitis; blood-retinal barrier dysfunction; adeno-associated virus; blindness; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A recombinant adeno-associated virus-vectored ribozyme composition,
                                                                                                                                                                                       09-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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                                                                                                                                                                                                                                                             ABZ72873 standard; RNA; 34 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                useful for treating a disease or dysfunction of the mammalian eye retinal disease, e.g. diabetic retinopathy or age-related macular
                                                                                                                                                 VEGFR2 Rz2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           exemplification
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     34 BP; 8
                                                                                                                                                                                                                                                                                                                                                                                                  AACUGAUGAGCCGUUCGCGGCGAAAC 30
                                                                                                                                               exemplary ribozyme construct
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           76.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         present invention
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Pred. No.
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                                                                                                                                               SEQ ID
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C first ribozyme that specifically cleaves an mRNA encoding a protein, and comprises (a) at rest a golypeptide, or peptide selected from the group of rod opsin, iNOS, CC RDS/peripherin, VEGFR1, VEGFR2, adenosine A-2B receptor, IGF-1, integrin alpha 1, integrin alpha 3, integrin alpha 5, or integrin alpha 4, or polymucleotide operably positioned downstream of at least a first CC promoter that directs expression of the polymucleotide in a selected CC mammalian cell transformed with the vector; (c) a viral particle CC comprising the ribozyme or the polymucleotide; or (e) a host cell comprising the ribozyme or the polymucleotide; or (e) a host cell CC comprising the ribozyme or the polymucleotide. Also described is a method CC for decreasing the amount of mRNA encoding a selected polypoptide in a cell composition described above, and for a time effective to specifically CC cleave the mRNA in the cell. (I) has ophthalmological activity, and can be used in gene therapy. (I) can be used for treating a disease or composition of the mammalian eye, such as a retinal disease or retinal degeneration. (I) is also useful for manufacturing a medicament for creating the diseases mentioned above, including autosomal dominant retinitis or a blood-retinal barrier dysfunction. (I) can also be useful for treating; decreasing the severity, or ameliorating the symptoms of a pathological condition, e.g. atrophic or pigmented lesions of the eye, but of the condition and condition, or a reduction in central or peripheral vision, or a reduction in the creating condition of the present invertion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A recombinant adeno-associated virus-vectored ribozyme composition, useful for treating a disease or dysfunction of the mammalian eye e retinal disease, e.g. diabetic retinopathy or age-related macular
                                         exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Page 81; 115pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     degeneration.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                describes a recombinant adeno-associated virus
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Sequence 34 BP; 10 A; 7 C; 9 G; 0 T; 8 U; 0 Other;

S Matches Query Match Best Local & u 26; Similarity **AACUGAUGAGCCGUUCGCGGCGAAAC** Conservative 100.0%; 76.5%; <u>.</u> Score 26; Pred. No. Mismatches DB 10; 0.00016 .00016; Length 34 0 Gaps

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AACUGAUGAGCCGUUCGCGGCGAAAC

ABZ72864 standard; RNA; 34

SXEXEXEX 09-APR-2003 (first entry)

ABZ72864;

Alpha V Rz2 exemplary ribozyme construct SEQ ID NO:99.

Hairpin ribozyme; hammerhead ribozyme; ribozyme; retinal disease; target;

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The present invention describes a recombinant adeno-associated virus (C (AAV) vectored ribozyme composition (I). (I) comprises: (a) at least a C first ribozyme that specifically cleaves an mRNA encoding a protein, C polypeptide, or peptide selected from the group of rod opsin, iNOS, C alpha 1, integrin alpha 3, integrin alpha 5, or integrin alpha 10; (b) a C vector comprising a polynucleotide encoding the ribozyme. Where the C polynucleotide operably positioned downstream of at least a first C promoter that directs expression of the polynucleotide in a selected C mammalian cell transformed with the vector; (c) a viral particle comprising the ribozyme or the polynucleotide; (d) an AAV vector C comprising the ribozyme or the polynucleotide; (d) an AAV vector C comprising the amount of mRNA encoding a selected polypeptide in a C retinal cell of a mammalian eye, comprising to the eye the C composition described above, and for a time effective to specifically C cleave the mRNA in the cell. (I) has ophthalmological activity, and can be used in gene therapy. (I) can be used for treating a disease or c treating the diseases mentioned above, including autosomal dominant C treating the diseases mentioned above, including autosomal dominant C retinitis or a blood-retinal barrier dysfunction. (I) can also be useful for treating the symptoms of a C pathological condition, e.g. atrophic or pigmented lesion, or a reduction in central or peripheral vision, or a reduction in central or peripheral vision.
                                                                                                         Query Match
Best Local S
Matches 26
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                                                                                                                                                                                                                                                          Sequence
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llarity 100.0%;
Conservative
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                                                                                                                                            Score 26; DB 10;
Pred. No. 0.00016;
                                                                                                             Mismatches
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The present invention describes a recombinant adeno-associated virus (C (AAV) vectored ribozyme composition (I) (I) comprises: (a) at least a CC first ribozyme that specifically cleaves an mRNA encoding a protein, CC polypeptide, or peptide selected from the group of rod opsin, 1NOS, CC RDS/peripherin, VEGFR1, VEGFR2, adenosine A-2B receptor, IGF-1, integrin CC alpha 1, integrin alpha 3, integrin alpha 5, or integrin alpha 4, integrin alpha 5, or integrin alpha 4, (b) a CC vector comprising a polymucleotide encoding the ribozyme, where the polymucleotide comparising the ribozyme or the polymucleotide; (c) a viral particle comprising the ribozyme or the polymucleotide; (d) an AAV vector comprising the ribozyme or the polymucleotide. Also described is a method CC comprising the amount of mRNA encoding a selected polypeptide in a critinal cell of a mammalian eye, comprising providing to the eye the CC composition described above, and for a time effective to specifically cleave the mRNA in the cell. (I) has ophthalmological activity, and can be used in gene therapy. (I) can be used for treating a disease or comprising the amount of mRNA encoding a selected macular codstinution of the mammalian eye, such as a retinal disease or retinal codstinution of the mammalian eye, such as a retinal disease or retinal codstinution of the mammalian eye, including autosomal dominant contraining the disease mantioned above, including autosomal dominant contraining the secretinal barrier dysfunction. (I) can also be useful for treating the disease or retinal dominant of the contraining the secretinal condition, e.g. atrophic or pipmented lesions of the eye, contraining the symptoms of a pathological condition, e.g. atrophic or pipmented lesions of the eye.
                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                          Sequence 34 BP; 10 A; 9 C; 10 G; 0 T; 5 U; 0 Other;
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CC blindness, a reduction in central or peripheral vision, or a reduction in total vision. ABZ72763 to ABZ72953 represent sequences used in the
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                                                                                     exemplification of the present invention
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                                                         C( AAV) vectored ribozyme composition (I). (I) comprises: (a) at least a cellipse of the specifically cleaves an mRNA encoding a protein, comprises: (a) at least a compriser, vector comprising selected from the group of rod opsin, iNOS, compriserin, VEGFR1, VEGFR2, adenosine A-2B receptor, IGP-1, integrin alpha 1, integrin alpha 3, integrin alpha 5, or integrin alpha V; (b) a comprising a polymucleotide encoding the ribozyme, where the comprising the ribozyme or the polymucleotide; (d) an AAV vector comprising the ribozyme or the polymucleotide; (d) an AAV vector comprising the ribozyme or the polymucleotide; (d) an AAV vector comprising the ribozyme or the polymucleotide; (d) an AAV vector comprising the ribozyme or the polymucleotide; (d) an AAV vector composition described above, and for a time effective to specifically composition described above, and for a time effective to specifically composition of the mammalian eye, comprising providing to the eye the composition of the mammalian eye, such as a retinal disease or composition of the mammalian eye, such as a retinal disease or composition of the mammalian eye, such as a retinal disease or retinal composition of the mammalian eye, such as a retinal disease or composition of the mammalian eye, such as a retinal disease or retinal composition of the disease and to a such composition of the mammalian eye, such as a retinal disease or composition of the disease and to a such composition of the disease or composition of the security or ameliorating the symptoms of a pathological condition, e.g. atrophic or pigmented lesions of the eye, composition or a reduction in central or peripheral vision, or a reduction in central or peripheral vision, or a reduction in central composition discount of the 
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Query Match Best Local : Matches

Similarity

73.5%;

Score 25; DB 10; Length 34; Pred. No. 0.00058;

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A recombinant adeno-associated virus-vectored ribozyme composition, useful for treating a disease or dysfunction of the mammalian eye e retinal disease, e.g. diabetic retinopathy or age-related macular e.g.

Lewin AS,

Shaw LC,

Grant MB

JAXD)

) UNIV FLORIDA

01-MAY-2001; 2001US-00847601

01-MAY-2002; 2002WO-US013679.

Claim 1; Page 80; 115pp; English.

CC (AAV) vectored ribozyme composition (I). (I) comprises: (a) at least a CC first ribozyme that specifically cleaves an mRNA encoding a protein, CC polypeptide, or peptide selected from the group of rod opsin, iNOS, CC RDS/peripherin, VEGFR1, VEGFR2, adenosine A-2B receptor, IGF-1, integrin alpha 3, integrin alpha 5, or integrin alpha V; (b) a CC vector comprising a polymuclacotide encoding the ribozyme, where the CC polymucleotide operably positioned downstream of at least a first CC promoter that directs expression of the polymucleotide in a selected CC mammalian cell transformed with the vector; (c) a viral particle CC comprising the ribozyme or the polymucleotide; (d) an AAV vector CC comprising the ribozyme or the polymucleotide; (d) an AAV vector CC comprising the ribozyme or the polymucleotide; or (e) a host cell CC comprising the ribozyme or the polymucleotide; or (e) a host cell CC comprising the mammalian eye, comprising providing to the eye the CC composition described above, and for a time effective to specifically CC cleave the mRNA in the cell. (I) has ophthalmological activity, and can be used in gene therapy. (I) can be used for treating a disease or retinal dysfunction, (diabetic) retinopathy, or (age-related) macular CC degeneration, (I) is also useful for manufacturing a medicament for treating the disease mentioned above, including autosomal dominant CC retinitis or a blood-retinal barrier dysfunction. (I) can also be useful for for the property or amedicating the symptoms of a composition of the manufacturing a medicament for the for the property or amedicating the symptoms of a composition of the manufacturing a medicament for the formation of decreating the second or composition or a blood-retinal barrier dysfunction. (I) can also be useful for the manufacturing a medicament for the formation of the manufacturing and the symptoms of a The present invention describes a recombinant adeno-associated virus for treating, decreasing the severity, or ameliorating the symptoms of a pathological condition, e.g. atrophic or pigmented lesions of the eye, blindness, a reduction in central or peripheral vision, or a reduction in

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RESULT 12
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ABZ712
CC (AAV) vectored ribozyme composition (I). (I) comprises: (a) at least a CC first ribozyme that specifically cleaves an mRNA encoding a protein, CC polypeptide, or peptide selected from the group of rod opsin, iNOS, CC RDS/peripherin, VEGFR1, VEGFR2, adenosine A-2B receptor, IGP-1, integrin alpha 1, integrin alpha 3, integrin alpha 5, or integrin alpha V; (b) a vector comprising a polynucleotide encoding the ribozyme, where the CC polynucleotide operably positioned downstream of at least a first CC promoter that directs expression of the polynucleotide in a selected CC mammalian cell transformed with the vector; (c) a viral particle CC comprising the ribozyme or the polynucleotide; (d) an AAV vector CC comprising the ribozyme or the polynucleotide; (e) a host cell CC comprising the ribozyme or the polynucleotide. Also described is a method CC for decreasing the amount of mRNA encoding a selected polypeptide in a CC cleave the mRNA in the cell. (I) has ophthalmological activity, and can CC be used in gene therapy. (I) can be used for treating a disease or cc dysfunction of the mammalian eye, such as a retinal disease or retinal CC dysfunction, (diabetic) retinopathy, or (age-related) macular for degeneration. (I) is also useful for manufacturing a medicament for
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes a recombinant adeno-associated virus (AAV) vectored ribozyme composition (I). (I) comprises: (a) at least
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    degeneration.
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CC (AAV) vectored ribozyme composition (1). (1) comprises: (a) at least a CC first ribozyme that specifically cleaves an mRNA encoding a protein, CC polypeptide, or peptide selected from the group of rod opsin, iNOS, CC RDS/peripherin, VEGFR1, VEGFR2, adenosine A-2B receptor, IGF-1, integrin alpha 1, integrin alpha 3, integrin alpha 5, or integrin alpha V; (b) a CC vector comprising a polymucleotide encoding the ribozyme, where the CC polymucleotide operably positioned downstream of at least a first CC promoter that directs expression of the polymucleotide in a selected CC comprising the ribozyme or the polymucleotide; (d) a viral particle CC comprising the ribozyme or the polymucleotide; (d) an AAV vector CC comprising the ribozyme or the polymucleotide, or (e) a host cell CC comprising the ribozyme or the polymucleotide. Also described is a method CC for decreasing the amount of mRNA encoding a selected polypeptide in a CC retinal cell of a mammalian eye, comprising providing to the eye the CC composition described above, and for a time effective to specifically
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A recombinant adeno-associated virus-vectored ribozyme composition, useful for treating a disease or dysfunction of the mammalian eye e retinal disease, e.g. diabetic retinopathy or age-related macular
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Best Local
(AAV) vectored ribozyme composition (I). (I) comprises: (a) at least first ribozyme that specifically cleaves an mRNA encoding a protein, polypeptide, or peptide selected from the group of rod opsin, iNOS, RDS/peripherin, VEGFRI, VEGFRI, adenosine A-2B receptor, IGF-1, integalpha 1, integrin alpha 3, integrin alpha 5, or integrin alpha V; (b) vector comprising a polynucleotide encoding the ribozyme, where the polynucleotide operably positioned downstream of at least a first promoter that directs expression of the polynucleotide in a selected mammalian cell transformed with the vector; (c) a viral particle comprising the ribozyme or the polynucleotide; (d) an AAV vector
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cleave the mRNA in the cell. (I) has ophthalmological activity, and can be used in gene therapy. (I) can be used for treating a disease or dysfunction of the mammalian eye, such as a retinal disease or retinal dysfunction, (diabetic) retinopathy, or (age-related) macular degeneration. (I) is also useful for manufacturing a medicament for treating the diseases mentioned above, including autosomal dominant retinities or a blood-retinal barrier dysfunction. (I) can also be useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A recombinant adeno-associated virus-vectored ribozyme composition useful for treating a disease or dysfunction of the mammalian eye retinal disease, e.g. diabetic retinopathy or age-related macular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hairpin ribozyme; hammerhead ribozyme; ribozyme; retinal disease; target; ophthalmological; gene therapy; eye; retinal dysfunction; AAV; diabetic retinopathy; macular degeneration; autosomal dominant retinitis; blood-retinal barrier dysfunction; adeno-associated virus; blindness; ss.
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                                                                                                                                                                                                                                                                                                  The present invention describes a recombinant adeno-associated virus (AAV) vectored ribozyme composition (I). (I) comprises: (a) at least
                                                                                                                                                                                                                                                                                                                                                                                                 Claim
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Best Local Similarity
Matches 25; Conserv
                                          The present invention relates to a method for identifying a gene with a selected function comprising contacting genes with a library of ribozymes and identifying at least 1 ribozyme that alters the selected function of the gene. The present sequence is an oligonucleotide used in the present invention. The methods (and ribozymes) are useful for identifying novel genes involved in retinal degradation, retinal disease, learning or memory, amylotropic lateral sclerosis or tumour suppression, and for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel methods for identifying genes with selected functions comprising contacting genes with a library of ribozymes, useful for identifying genes involved in, e.g. retinal disease, learning or memory and tumor
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producing non-human animal models of diseases

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                                                                     Sequence 44 BP; 13 A; 11 C; 10 G; 10 T; 0 U; 0 Other;
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                6 ACUGAUGAGCCGUUCGCGGCGAAAC 30
                                            Similarity
                                  Conservative
                                           73.5%;
84.0%;
                                 4
                                          Score 25;
Pred. No.
                                  Mismatches
35
                                           0.00057;
                                                     DB 3;
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Search completed: August 28, 2006, 09:50:06 Job time : 286 scas

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Maximum DB
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and is derived by analysis of the total score distribution
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/EMC_Celerra_SIDS3/ptodata/2/pubpna/US08_PUBCOMB.seq:*
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Copyright (c) 1993 - 2006 Biocceleration Ltd
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US-09-847-601B-91
US-09-847-601B-96
US-09-847-601B-97
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                                                                               Sequence 100, App
Sequence 98, Appl
Sequence 92, Appl
Sequence 93, Appl
Sequence 99, Appl
Sequence 90, Appl
Sequence 91, Appl
Sequence 91, Appl
Sequence 91, Appl
Sequence 95, Appl
Sequence 97, Appl
Sequence 91, Appl
Sequence 91, Appl
Sequence 91, Appl
Sequence 92, Appl
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| 5                    | 44              | 43                 | 42              | 41              | 40              | 39                  | 38                   | 37               | 36                 | კ<br>5               | 34<br>4              | 33                 | 32                   | 31                 | 30                    | 29                    | 28                    | 27                    | 26       | 25                   | 24                    | 23                    | 22                 | 21       | 20                  | 19                  | 18                 |
| 15                   | 15              | 15                 | 15              | 15              | 15              | 15                  | 15                   | 15               | 15                 | 15                   | 15                   | 15                 | 15                   | 15                 | 15                    | 15                    | 15                    | 15                    | 15       | 15                   | 15                    | 15                    | 15                 | 15       | 15                  | 15                  | 15                 |
| 44.1                 | 44.1            | 44.1               | 44.1            | 44.1            | 44.1            | 44.1                | 44.1                 | 44.1             | 44.1               | 44.1                 | 44.1                 | 44.1               | 44.1                 | 44.1               | 44.1                  | 44.1                  | 44.1                  | 44.1                  | 44.1     | 44.1                 | 44.1                  | 44.1                  | 44.1               | 44.1     | 44.1                | 44.1                | 44.1               |
| 1413                 | 1413            | 1413               | 1401            | 1155            | 1155            | 974                 | 957                  | 954              | 939                | 873                  | 852                  | 810                | 714                  | 634                | 575                   | 575                   | 575                   | 575                   | 573      | 522                  | 389                   | 389                   | 379                | 343      | 307                 | 250                 | 200                |
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| US-10-282-122A-20379 | -10-369-493-245 | US-09-815-242-6074 | US-10-181-906-5 | US-10-126-764-9 | US-10-309-515-9 | US-10-198-846-13796 | US-10-282-122A-28269 | US-10-493-462-40 | US-10-156-761-4428 | US-10-282-122A-12733 | US-10-282-122A-26469 | US-10-156-761-2123 | US-10-282-122A-14226 | US-10-094-097B-106 | US-09-925-065A-637542 | US-09-925-065A-637541 | US-09-925-065A-637542 | US-09-925-065A-637541 | 1        | US-10-282-122A-11370 | US-09-925-065A-554407 | US-09-925-065A-554407 | US-09-867-701-4078 | -10      | US-10-437-963-99234 | US-09-864-761-31920 | US-09-960-352-4463 |
| Sequence             | Seguence        | Sequence           | Sequence        | Sequence        | Sequence        | Sequence            | Sequence             | Sequence         |                    |                      | Sequence             | Sequence           | Sequence             | Sequence           |                       |                       |                       | Sequence              | Sequence | Sequence             | Sequence              | Sequence              | Sequence           | Sequence | Sequence            | Sequence            | Sequence           |
| 20379, A             |                 | 6074, Ap           | 5, Appli        | 9, Appli        | 9, Appli        | 13796, A            | 28269, A             | 40, Appl         | 4428, Ap           | 12733, A             | 26469, A             | 2123, Ap           | 14226, A             | 106, App           | 637542,               | 637541,               | 637542,               | 637541,               | 15387, A | 11370, A             | 554407,               | 554407,               | 4078, Ap           | 36422, A | 99234, A            | 31920, A            | 4463, Ap           |

## ALIGNMENTS

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GENERAL INFORMATION:

APPLICANT: LEWIN, ALFRED S.

APPLICANT: SANW, LYNK C.

APPLICANT: GRAM, LYNK C.

APPLICANT: GRAM, MARIA B.

TITLE OF INVENTION: METHODS FOR THE TREATMENT OF RETINAL DISEASES

FILE REFERENCE: 4300.014100

CURRENT APPLICATION NUMBER: US/09/847,601B

CURRENT FILING DATE: 2001-05-01

PRIOR APPLICATION NUMBER: 09/063,667

PRIOR APPLICATION NUMBER: 09/063,667

PRIOR FILING DATE: 1998-04-21

PRIOR APPLICATION NUMBER: 60/046,147

PRIOR APPLICATION NUMBER: 60/046,147
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US-09-847-601B-100
                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 182
SOFTWARE: PatentIn version 3.2
SEQ ID NO 100
                                                                                                      Matches
                                                                                                                            Best
                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 100, Appropriate Publication No.
                                                                                                                                                                                                                                                            TYPE: RNA
ORGANISM: Artificial
FEATURE:
                                                                                                                                                                                                                                   OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                          LENGTH: 34
                                                                                                                            Local
                                1 UGCAAACUGAUGAGCCGUUCGCGGCGAAACGAAG 34
                                                                                                 h 100.0%; Score 34; DI
Similarity 100.0%; Pred. No. 3.7
34; Conservative 0; Mismatches
UGCANACUGAUGAGCCGUUCGCGGCGAAACGAAG
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o. US20050096282A1
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RESULT 2 US-09-847-601B-101

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NUMBER OF SEQ ID NOS: 182
SOFTWARE: Patentin version 3.2
SEQ ID NO 98
LENGTH: 34
TYPE: RNA
ORGANISM: Artificial
FEATURE:
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                                                                                                                                                                                           ; OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE US-09-847-601B-98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE US-09-847-601B-101
                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT FILING DATE: 2001-05-01
PRIOR APPLICATION NUMBER: 09/063,667
PRIOR FILING DATE: 1998-04-21
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR PILING DATE: 1997-05-09
PRIOR PPLICATION NUMBER: 60/044,492
PRIOR PILING DATE: 1997-04-21
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SEQ ID NO 101
LENGTH: 32
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APPLICANT: SHAW, LYNN C.
APPLICANT: GRANT, MARIA B.
TITLE OF INVENTION: ADENO-ASSOCIATED VIRUS-DELIVERED RIBOZYME COMPOSITIONS AND TITLE OF INVENTION: METHODS FOR THE TREATMENT OF RETINAL DISEASES
FILE REFERENCE: 4300.014100
CURRENT APPLICATION NUMBER: US/09/847,601B
CURRENT FILING DATE: 2001-05-01
PRIOR APPLICATION NUMBER: 09/063,667
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR PILING DATE: 1997-05-09
PRIOR APPLICATION NUMBER: 60/046,4492
PRIOR FILING DATE: 1997-04-21
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PRIOR PRIOR PRIOR DATE: 1997-04-21
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APPLICANT: SHAW, LYNN C.
APPLICANT: GRANT, MARIA B.
APPLICANT: GRANT, MARIA B.
APPLICANT: GRANT, MARIA B.
TITLE OF INVENTION: ADENO-ASSOCIATED VIRUS-DELIVERED RIBOZYME COMPOSITIONS AND TITLE OF INVENTION: METHODS FOR THE TREATMENT OF RETINAL DISEASES
FILE REFERENCE: 4300.014100
CURRENT APPLICATION NUMBER: US/99/847,601B
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ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                                                             Local Similarity hes 27; Conserv
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                                              AACUGAUGAGCCGUUCGCGGCGAAACG 31
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o. US20050096282A1
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                                                                                                                    79.4%;
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                                                                                             Mismatches
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APPLICANT: LEWIN, ALFRED S.
APPLICANT: SHAW, LYNN C.
APPLICANT: SHAW, LYNN C.
APPLICANT: GRANT, MARIA B.
TITLE OF INVENTION: ADENO-ASSOCIATED VIRUS-DELIVERED RIBOZYME COMPOSITIONS AND TITLE OF INVENTION: METHODS FOR THE TREATMENT OF RETINAL DISEASES
FILE REFERENCE: 4300.014100
CURRENT APPLICATION NUMBER: US/09/847,601B
CURRENT FILING DATE: 2001-05-01
FRIOR APPLICATION NUMBER: 60/063,667
PRIOR APPLICATION NUMBER: 60/066,147
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR FILING DATE: 1997-05-09
PRIOR APPLICATION NUMBER: 60/044,492
PRIOR FILING DATE: 1997-05-09
PRIOR FILING DATE: 1997-05-09
FRIOR FILING DATE: 1997-05-09
PRIOR APPLICATION NUMBER: 60/044,492
PRIOR FILING DATE: 1997-05-09
PRIOR APPLICATION NUMBER: 60/044,492
PRIOR FILING DATE: 1997-05-09
PRIOR APPLICATION NUMBER: 60/044,492
PRIOR FILING DATE: 1997-05-09
PRIOR PRI
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SOFTWARE: Patentin version 3.2
SEQ ID NO 92
LENGTH: 34
TYPE: RNA
ORGANISM: Artificial
FEATURE:
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US-09-847-601B-93
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Matches
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                                    Best Local
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CURRENT APPLICATION NUMBER: US/09/847,601B
CURRENT FILING DATE: 2001-05-01
PRIOR APPLICATION NUMBER: 09/063,667
PRIOR FILING DATE: 1998-04-21
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR FILING DATE: 1997-05-09
PRIOR FILING DATE: 1997-05-09
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: ADENO-ASSOCIATED VIRUS-DELIVERED RIBOZYME COMPOSITIONS AND TITLE OF INVENTION: METHODS FOR THE TREATMENT OF RETINAL DISEASES
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                                                                                                                                                                                     FEATURE:
OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE
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Similarity 100.0%; Pred. No. 2.3e-(
26; Conservative 0; Mismatches
                                    Similarity
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Conservative
                                100.0%;
                                76.5%; Score 26; 100.0%; Pred. No.
Mismatches
                                DB 4; I
2.3e-05;
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                                                                                                                                                                                                                           SOFTWARE: PatentIn version 3.2 SEQ ID NO 90
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Best Local Similarity
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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                     APPLICANT: LEWIN, ALFRED S.
APPLICANT: SHAW, LYNN C.
APPLICANT: GRAWT, MARIA B.
TITLE OF INVENTION: ADENO-ASSOCIATED VIRUS-DELIVERED RIBOZYME COMPOSITIONS AND FITTE OF INVENTION: METHODS FOR THE TREATMENT OF RETINAL DISEASES FILE REFERENCE: 4300.014100
CURRENT APPLICATION NUMBER: US/09/847,601B
CURRENT FILING DATE: 2001-05-01
PRIOR APPLICATION NUMBER: 09/063,667
PRIOR APPLICATION NUMBER: 60/046,147
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PRIOR FILING DATE: 1997-05-09
PRIOR APPLICATION NUMBER: 60/044,492
PRIOR FILING DATE: 1997-04-21
NUMBER OF SEQ ID NOS: 182
SOFTWARE: PatentIn version 3.2
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CURRENT FILING DATE: 2001-05-01
PRIOR APPLICATION NUMBER: 09/063,667
PRIOR FILING DATE: 1998-04-21
                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 60/044,492
PRIOR FILING DATE: 1997-04-21
NUMBER OF SEQ ID NOS: 182
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APPLICANT: SHAW, LYNN C.
APPLICANT: GRANT, MARIA B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 34
TYPE: RNA
ORGANISM: Artificial
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                                                                                                                         ORGANISM: Artificial FEATURE:
                                                                                                                                                                           TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
                                                                                                    OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE
                                                                                                                                                                                                        ENGTH: 34
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73.5%;
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100.0%; Pred. No. 2.3e-05;
ive 0; Mismatches 0;
Score 25; DB 4; Length 34; Pred. No. 9.1e-05;
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OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE US-09-847-601B-95
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SEQ ID NO 91
LENGTH: 34
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                                                                                                          SOFTWARE: PatentIn version 3.2
SEQ ID NO 95
LENGTH: 34
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                                                                                                                                                                                       APPLICANT: LEWIN, ALFRED S.
APPLICANT: SHAW, LYNN C.
APPLICANT: GRAWT, MARIA B.
APPLICANT: GRAWT, MARIA B.
TITLE OF INVENTION: ADEND-ASSOCIATED VIRUS-DELIVERED RIBOZYME COMPOSITIONS AND TITLE OF INVENTION: METHODS FOR THE TREATMENT OF RETINAL DISEASES
FILE REFERENCE: 4300.014100

CURRENT FILLING DATE: 2001-05-01
PRIOR APPLICATION NUMBER: 09/063,667
PRIOR PILLING DATE: 1998-04-21
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR APPLICATION NUMBER: 60/044,492
PRIOR PILLING DATE: 1997-05-09
PRIOR PILLING DATE: 1997-04-21
PRIOR PILLING DATE: 1997-04-21
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APPLICANT: SHAW, LYNN C.
APPLICANT: GRANT, MARIA B.

APPLICANT: GRANT, MARIA B.

TITLE OF INVENTION: ADENO-ASSOCIATED VIRUS-DELIVERED RIBOZYME COMPOSITIONS AND TITLE OF INVENTION: METHODS FOR THE TREATMENT OF RETINAL DISEASES FILE REFERENCE: 4300.014100

CURRENT FILLING DATE: 2001-05-01

PRIOR APPLICATION NUMBER: 09/063,667

PRIOR APPLICATION NUMBER: 60/046,147

PRIOR APPLICATION NUMBER: 60/046,147

PRIOR APPLICATION NUMBER: 60/044,492

PRIOR PILING DATE: 1997-05-09

PRIOR PILING DATE: 1997-04-21

PRIOR PILING DATE: 1997-04-21

PRIOR PILING DATE: 1997-04-21

PRIOR PILING DATE: 1997-04-21
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                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 182
                                                                 ORGANISM: Artificial
                                                                                           TYPE: RNA
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ORGANISM: Artificial
                                             FEATURE:
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100.0%; Pr
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100.0%; Pred. No. 9.
Live 0; Mismatches
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Sequence 97, Application US/09847601B
Publication No. US20050096282A1
GENERAL INFORMATION:
APPLICANT: LEWIN, ALFRED S.
APPLICANT: SHAW, LYNN C.
APPLICANT: GRANT, MARIA B.
TITLE OF INVENTION: METHODS FOR THE TREATMENT OF RETINAL DISEASES
TITLE OF INVENTION UNMERS: US/09/847,601B
CURRENT APPLICATION NUMBER: US/09/847,601B
CURRENT FILING DATE: 1998-04-21
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR FILING DATE: 1997-05-09
PRIOR APPLICATION NUMBER: 60/044,492
PRIOR FILING DATE: 1997-09-21
NUMBER OF SEQ ID NOS: 182
SOFTWARE: Patentin version 3.2
FRIOR PATENTIN VERSION 3.2
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Best Local Similarity
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Best Local Similarity
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CURRENT PEDLICATION NUMBER: US/09/847,601B

CURRENT FILING DATE: 2001-05-01

PRIOR APPLICATION NUMBER: 09/063,667

PRIOR PILING DATE: 1998-04-21

PRIOR PILING DATE: 1997-05-09

PRIOR APPLICATION NUMBER: 60/046,147

PRIOR PILING DATE: 1997-05-09

PRIOR PILING DATE: 1997-05-09

PRIOR PILING DATE: 1997-04-21

NUMBER OF SEQ ID NOS: 182

SOFTWARE: Patentin version 3.2
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APPLICANT: SHAW, LYNN C.
APPLICANT: GRANT, MARIA B.
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ORGANISM: Artificial
FEATURE:
TYPE: RNA ORGANISM: Artificial
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Pred. No. 9.1e-05,
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Pred. No. 9.1e-05;
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APPLICANT: LEWIN, ALFRED S.
APPLICANT: SHAW, LYNN C.
APPLICANT: SHAW, LYNN C.
APPLICANT: GEANT, MARIA B.
TITLE OF INVENTION: ADENO-ASSOCIATED VIRUS-DELIVERED RIBOZYME COMPOSITIONS AND TITLE OF INVENTION: METHODS FOR THE TREATMENT OF RETINAL DISEASES
FILE REFERENCE: 4300.014100
CURRENT APPLICATION NUMBER: US/09/847,601B
CURRENT FILING DATE: 2001-05-01
PRIOR APPLICATION NUMBER: 09/63,667
PRIOR FILING DATE: 1998-04-21
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR APPLICATION NUMBER: 60/046,147
PRIOR FILING DATE: 1997-05-09
PRIOR APPLICATION NUMBER: 60/044,492
PRIOR APPLICATION NUMBER: 60/044,492
PRIOR FILING DATE: 1997-04-21
NUMBER OF SEQ ID NOS: 182
SOFTWARE: Patentin version 3.2
SEQ ID NO 94
LENGTH: 34
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; OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE
US-09-847-601B-97
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SEQ ID NO 3252
LENGTH: 463
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Publication No. US20040123343A1
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Best Local Similarity
                                                                       APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
                                                                                                                                                                                                                                                                APPLICANT: La Rosa, Thomas J. APPLICANT: Kovalic, David K. APPLICANT: Zhou, Yihua
                                         CURRENT FILING DATE:
NUMBER OF SEQ ID NOS:
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Wu, Wei
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No. US20050096282A1
                                                                                                                                                                              Boukharov, Andrey A.
Barbazuk, Brad
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Pred. No.
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9.1e-05;
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; TYPE: DNA; ORGANISM: Streptomyces avermitilis; FRATURE: ; RAME/KEY: CDS; LOCATION: (1)...(1845)
US-10-156-761-3619
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; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT4530_102953C.1
US-10-437-963-3252
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APPLICANT: SAKAKI, YOSHIYUKI
APPLICANT: HATTORI, MASAHIRA
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES
FILE REFERENCE: 249-262
FILE REPERICATION NUMBER: US/10/156,761
CURRENT APPLICATION NUMBER: J072-05-29
PRIOR APPLICATION NUMBER: JD 2001-204089
PRIOR FILING DATE: 2001-05-30
PRIOR FILING DATE: 2001-05-30
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Best Local Similarity
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PRIOR FILING DATE: 2001-08-02
NUMBER OF SEQ ID NOS: 15109
SEQ ID NO 3619
SEQ ID NO 18859
                   APPLICANT: La Roba, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Wu, Wei
APPLICANT: Wu, Wei
APPLICANT: Boukharov, Andrey A.
APPLICANT: Barbazuk, Brad
APPLICANT: Li, Ping
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
TITLE OF INVENTION: Which Plants and Uses Thereof for Plant Improvement
TITLE OF INVENTION: WISHELD BARBASH BARBA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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Best Local Similarity
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APPLICANT: IKEDA, HARUO
APPLICANT: ISHIKAWA, JUN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ENGTH: 1845
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221 TGCAAACTGATGAGCC 206
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HORIKAWA, HIROSHI
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81.2%;
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Pred. No. 22;
3; Mismatches
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Search completed: August 28, 2006, 10:41:09 Job time : 718 secs

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US-11-266-748A-25189

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| .2 777 8 US-11-266-748A-174235 Sequence 172 777 8 US-11-266-748A-190325 Sequence 182 778 9 US-11-266-748A-190325 Sequence 182 820 8 US-11-266-748A-189268 Sequence 182 918 8 US-11-266-748A-352337 Sequence 182 918 8 US-11-266-748A-35736 Sequence 352 918 8 US-11-266-748A-35736 Sequence 352 1169 6 US-10-520-999-1 Sequence 42 1210 6 US-10-520-999-1 Sequence 42 1229 6 US-10-520-999-3 Sequence 22 1280 6 US-10-520-999-3 Sequence 32 1280 6 US-10-520-999-3 Sequence 22 1280 6 US-10-520-999-3 Sequence 22 1280 6 US-10-520-999-3 Sequence 32 1607 6 US-10-520-999-3 Sequence 32 1607 6 US-10-520-999-3 Sequence 32 1607 6 US-10-520-999-3 Sequence 42 2 1812 6 US-11-218-305-10001 Sequence 42 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2   |   | 14   | 14   | 14                | 14   | 14     | 14   | 14   | 14     | 14    | 14   | 14   | 14   | 14              | 14   | 14       | 14   | 14  | 14  | 14  | 14     | 14  | 14                 |
| B US-11-266-748A-174235 Sequence 179 US-11-266-748A-190325 Sequence 189 US-11-266-748A-189268 Sequence 189 US-11-266-748A-35237 Sequence 189 US-11-266-748A-35716 Sequence 189 US-11-266-748A-35716 Sequence 189 US-11-266-748A-35716 Sequence 110 US-10-520-999-1 Sequence 110 US-10-520-999-1 Sequence 2, US-10-520-999-2 Sequence 3, US-10-520-999-2 Sequence 2, US-10-520-999-2 Sequence 100 US-10-471-571A-4927 Sequence 110 US-10-520-999-6 Sequence 110 US-11-218-305-10001 Sequence 130 US-11-218-305-1002 Sequence 130 US-11-218-305-1002 Sequence 130 US-11-266-748A-24493 Sequence 238 US-11-266-748A-24493 Sequence 238 US-11-266-748A-28460 Sequence 289  |   | 41.2 |      |                   |      |        |      |      |        |       |      |      |      |                 |      |          |      |     |     |     |        |     |                    |
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| 1797<br>1797<br>1797<br>1797<br>1797<br>1797<br>1797<br>1797   |   | J.   | i    | -11-266-748A-2449 | .1   | Ĭ.     | ï    | ١.   |        | Ļ     | .1.  | 1    |      | US-10-520-999-1 | •    | ŀ        |      | ï   | H   | -11 | -11    | 11  | -11-266-748A-17423 |
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## ALIGNMENTS

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APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: Bio-oriented Technology Research Advancement Institution
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
ITITLE OF INVENTION: FULL-LENGTH PLANT cDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
US-11-266-748A-184169
; Sequence 184169, Application US/11266748A
; Publication No. US20060134663A1
; GENERAL INFORMATION:
; APPLICANT: Harkin, Paul
; APPLICANT: Johnston, Patrick
; APPLICANT: Mulligan, Karl
; TITLE OF INVENTION: Transcriptome Microarray Technology and
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US-10-449-902-27861
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 27861
LENGTH: 1809
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PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: AK103303
DATABASE ENTRY DATE: 2002-08-28
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                                                                                                                                                                                                                                                                                                                     12 GAGCCGUUCGCGGCGA 27
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537 GAGCCGTTCGCGGCGA 552
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Similarity 87.5%; Pred. No. 2.7
14; Conservative 2; Mismatches
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E OF INVENTION: Methods of Using the Same REFERENCE: 55815-0102 (319189)

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; TYPE: DNA
; ORGANISM: Homo Sapiens
US-11-266-748A-191642
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 191642, Application US Publication No. US20060134663A1 GENERAL INFORMATION:
                                                                                     SOFTWARE: PatentIn
SEQ ID NO 191642
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LENGTH: 1084
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                                                                                                                                                      PRIOR APPLICATION NUMBER: US 60/700,293 PRIOR FILING DATE: 2005-07-18
                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT FILING DATE: 2005-11-03
PRIOR APPLICATION NUMBER: EP 04105479.2
PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105482.6
PRIOR FILING DATE: 2004-11-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/11/266,748A CURRENT FILING DATE: 2005-11-03 PRIOR APPLICATION NUMBER: EP 04105479.2 PRIOR FILING DATE: 2004-11-03 PRIOR APPLICATION NUMBER: EP 04105482.6
                                                                                                                                  NUMBER OF SEQ ID NOS: 483996
                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105507.0
                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: EP 04105483.4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: 55815-0102 (319189)
CURRENT APPLICATION NUMBER: US/11/266,748A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Transcriptome Microarray Technology and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Harkin,
APPLICANT: Johnst
APPLICANT: Mullig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn
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PRIOR FILING DATE: 2005-07-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: EP 04105483.4 PRIOR FILING DATE: 2004-11-03
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                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 2004-11-03
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TYPE: DNA
ORGANISM: Homo Sapiens
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                                                                 LENGTH:
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                                                                                                                                                                                                APPLICATION NUMBER: EP 04105484.2 FILING DATE: 2004-11-03 APPLICATION NUMBER: US 60/662,276 FILING DATE: 2005-03-14
                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: EP 04105485.9 FILING DATE: 2004-11-03
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APPLICATION NUMBER: US 60/662,276
FILING DATE: 2005-03-14
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Mulligan, Karl
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73.3%;
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TITLE OF INVENTION: Transcriptome Microarray Technology and TITLE OF INVENTION: Methods of Using the Same FILE REFERENCE: 55815-0102 (319189)
CURRENT APPLICATION NUMBER: US/11/266,748A
CURRENT FILING DATE: 2005-11-03
PRIOR APPLICATION NUMBER: EP 04105479.2
PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105482.6
PRIOR FILING DATE: 2004-11-03
PRIOR PILING DATE: 2004-11-03
PRIOR FILING DATE: 2004-11-03
PRIOR FILING DATE: 2004-11-03
PRIOR PILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105403.4
PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105405.9
PRIOR APPLICATION NUMBER: EP 04105405.9
PRIOR PILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105405.9
PRIOR PILING DATE: 2004-11-03
PRIOR PILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105405.9
PRIOR PILING DATE: 2004-11-03
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US-10-449-902-20318
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US-11-266-748A-25135
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Best Local (
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TITLE OF INVENTION: FULL-LENGTH PLANT CDNA FILE REFERENCE: MOA-A0205Y1-US CURRENT APPLICATION NUMBER: US/10/449,902 CURRENT FILING DATE: 2003-05-29 PRIOR APPLICATION NUMBER: JP 2002-203269 PRIOR FILING DATE: 2002-05-30 PRIOR PILING DATE: 2002-12-11
                                                                                                                                                                                  APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: Bio-oriented Technology Research Advancement Ins
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Scie
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PRIOR FILING DATE: 2005-03-14
PRIOR APPLICATION NUMBER: US 60/700,293
PRIOR FILING DATE: 2005-07-18
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APPLICANT: Mulligan, Karl
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ORGANISM: Homo Sapiens
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Pred. No.
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Pred. No.
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US-11-266-748A-251890/c
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Publication No. US20060123505A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Harkin, Paul
                                                                                                                                                                                                                                                                                  Sequence 251890, Application US/11266748A Publication No. US20060134663A1
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 4455
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Best Local
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 20318
LENGTH: 1243
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            APPLICANT: Johnston, Patrick
APPLICANT: Mulligan, Karl
TITLE OP INVENTION: Transcriptome Microarray Technology and
TITLE OP INVENTION: Methods of Using the Same
FILE REFERENCE: 55815-0102 (319189)
CURRENT APPLICATION NUMBER: US/11/266,748A
CURRENT APPLICATION NUMBER: US/11/266,748A
CURRENT FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105479.2
PRIOR FILING DATE: 2004-11-03
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CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL LENGTH PLANT CDNA AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: OTYZA BALIVA
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: AK061781
DATABASE ENTRY DATE: 2001-12-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: OTYZA BATIVA
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: AK099148
DATABASE ENTRY DATE: 2002-08-28
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RIOR APPLICATION NUMBER: EP 04105482.6
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405 GCAAACTGATGAGCC 419
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Pred. No. 11;
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; ORGANISM: Homo Sapiens
US-11-266-748A-312407
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PRIOR FILING DATE: 2004-11-03
PRIOR PELICATION NUMBER: EP 04105482.6
PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105483.4
PRIOR FILING DATE: 2004-11-03
PRIOR FILING DATE: 2004-11-03
PRIOR FILING DATE: 2004-11-03
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PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105484.2
PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: US 60/662,276
PRIOR APPLICATION NUMBER: US 60/700,293
PRIOR APPLICATION NUMBER: US 60/700,293
PRIOR PILING DATE: 2005-07-18
                                                            Matches
                                                                                                Query Match
                                                                                                                                                                                                                        SEQ ID NO 312407
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PRIOR FILING DATE: 2005-03-14
PRIOR APPLICATION NUMBER: US 60/700,293
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CURRENT APPLICATION NUMBER: US/11/266,748A
CURRENT FILING DATE: 2005-11-03
                                                                                                                                                                                                                                               SOFTWARE: PatentIn
                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 483996
                                                                                                                                                                                                                                                                                       PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105484.2
PRIOR FILING DATE: 2004-11-03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Johnston, Patr
APPLICANT: Mulligan, Karl
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NUMBER OF SEQ ID NOS: 483996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
                                                                                                                                                                                    TYPE: DNA
                                                                                                                                                                                                       ENGTH: 1342
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APPLICATION NUMBER: EP 04105483.4
FILING DATE: 2004-11-03
APPLICATION NUMBER: EP 04105507.0
FILING DATE: 2004-11-03
                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: EP 04105485.9
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6 ACUGAUGAGCCGUUC
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                                                                                 Similarity
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                                                            Conservative
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                                                                             44.18;
73.38;
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                                                                               Score 15; DE
Pred. No. 11;
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Pred. No. 11;
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                                                            Mismatches
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                                                               Indels
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1025 ACTGATGAGCCGTTC 1039

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; OTHER INFORMATION: Pratylenchus scribneri rDNA
US-11-256-428-61
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US-11-256-428-61/c
                    US-11-266-748A-184170
                                        RESULT 11
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Sequence 184170, Application US/11266748A
                                                                                                                                                                              Matches
                                                                                                                                                                                               Query Match
Best Local
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SEQ ID NO 61
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SEQ ID NO 20725
LENGTH: 1551
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                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Niblett, Charles L.
TITLE OF INVENTION: Methods and Materials for Conferring Resistance to Pests
TITLE OF INVENTION: Pathogens of Plants
FILE REFERENCE: VEN-100
CURRENT APPLICATION NUMBER: US/11/256,428
CURRENT FILING DATE: 2005-10-21
NUMBER OF SEQ ID NOS: 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: JF 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: OTYZA BATIVA
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: AK071031
DATABASE ENTRY DATE: 2001-12-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
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                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                       LENGTH: 1773
TYPE: DNA
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12; Conserv
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86.78;
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80.0%;
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Pred. No.
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Pred. No. 11;
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US-11-266-748A-191643
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                  FILING DATE:
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PRIOR APPLICATION NUMBER: EP 04105479.2
PRIOR PILING DATE: 2004-11-03
PRIOR PILING DATE: 2005-03-14
PRIOR PILING DATE: 2005-03-14
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                                                                                                                                                                                                                                                                            APPLICANT: Harkin, Paul
APPLICANT: Johnston, Patrick
APPLICANT: Mulligan, Karl
TITLE OF INVENTION: Transcriptome Microarray Technology and
TITLE OF INVENTION: Methods of Using the Same
FILE REFERENCE: 55815-0102 (319189)
CURRENT APPLICATION NUMBER: US/11/266,748A
CURRENT FILING DATE: 2005-11-03
CURRENT FILING DATE: 2005-11-03
PRIOR APPLICATION NUMBER: EP 04105479.2
                                                                                                          PRIOR APPLICATION NUMBER: EP 04105483.4
PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105507.0
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PRIOR APPLICATION NUMBER: EP 04105482.6
PRIOR FILING DATE: 2004-11-03
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CURRENT APPLICATION NUMBER: US/11/266,748A
CURRENT FILING DATE: 2005-11-03
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APPLICANT: Johnston, Patr
APPLICANT: Mulligan, Karl
                  PRIOR FILING DATE: 2004-11-03
PRIOR APPLICATION NUMBER: EP 04105485.9
PRIOR FILING DATE: 2004-11-03
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LOCATION: (2038)..(2071)
OTHER INFORMATION: n is a,
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TYPE: DNA
ORGANISM: Homo Sapiens
FEATURE:
APPLICATION NUMBER: EP 04105484.2
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Similarity 73.3%;
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Pred. No.
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Query Match
Best Local Similarity
warches 11; Conserve
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; TYPB: DNA
; ORGANISM: Homo sapiens
US-11-226-554-73
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; NAME/KEY: misc feature
; LOCATION: (2038)..(2071)
; OTHER INFORMATION: n is a, c,
US-11-266-748A-191643
          S-11-248-718-73
Sequence 73, Application US/11248718
Publication No. US20060160997A1
GENERAL INFORMATION:
APPLICANT: Cairns, Belinda
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PRIOR APPLICATION NUMBER: US 60/700,293
PRIOR FILING DATE: 2005-07-18
NUMBER OF SEQ ID NOS: 483996
SOFTWARE: Patentin version 3.3
SEQ ID NO 191643
LENGTH: 2253
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CURRENT FILING DATE: 2005-09-13
FRIOR APPLICATION NUMBER: US 10/177,488
PRIOR FILING DATE: 2002-06-19
PRIOR APPLICATION NUMBER: US 60/366,869
PRIOR FILING DATE: 2002-03-20
PRIOR FILING DATE: 2002-03-20
NUMBER OF SEQ ID NOS: 154
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: P5001R1C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Compositions and Methods TITLE OF INVENTION: Treatment of Tumor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
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                                                                                                                                             746 ACTGATGAGCCGTTC 760
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Phillips, Heidi S.
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Hillan, Kenneth J.
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                                                                                                                                                                                                             Conservative
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73.38;
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Pred. No. 11;
4; Mismatches
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Pred. No. 11;
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TITLE OF INVENTION: Transcriptome Microarray Technology and TITLE OF INVENTION: Methods of Using the Same FILE REFERENCE: 55815-0102 (319189)
CURRENT APPLICATION NUMBER: US/11/266,748A
CURRENT APPLICATION NUMBER: US/11/266,748A
CURRENT FILING DATE: 2005-11-03
PRIOR PILLING DATE: 2004-11-03
PRIOR FILLING DATE: 2004-11-03
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PRIOR APPLICATION NUMBER: EP 04105403.4
PRIOR APPLICATION NUMBER: BD 04105507.0
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US-11-266-748A-29125
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; ORGANISM: Homo Sapien
US-11-248-718-73
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CURRENT APPLICATION NUMBER: US/11/248,718
CURRENT TILING DATE: 2005-10-11
PRIOR APPLICATION NUMBER: US/10/177,488
PRIOR FILING DATE: 2002-06-19
PRIOR APPLICATION NUMBER: US 60/299,500
PRIOR FILING DATE: 2001-06-20
PRIOR PILING DATE: 2001-06-20
PRIOR APPLICATION NUMBER: US 60/300,880
PRIOR APPLICATION NUMBER: US 60/301,880
PRIOR FILING DATE: 2001-06-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 29125, Application US/11266748A Publication No. US20060134663A1 GENERAL INFORMATION:
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Best Local Similarity
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APPLICANT: Johnston, Patrick
APPLICANT: Mulligan, Karl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 2001-11-07
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 154
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TITLE OF INVENTION: Compositions and Methods for the Diagnosis and
TITLE OF INVENTION: Treatment of Tumor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OR APPLICATION NUMBER: US 60/304,813
OR FILING DATE: 2001-07-11
OR APPLICATION NUMBER: US 60/312,312
OR FILING DATE: 2001-08-13
OR APPLICATION NUMBER: US 60/314,280
OR APPLICATION NUMBER: US 60/314,280
OR PILING DATE: 2001-08-22
OR APPLICATION NUMBER: US 60/339,227
OR APPLICATION NUMBER: US 60/339,227
OR APPLICATION NUMBER: US 60/339,227
OR FILING DATE: 2001-10-19
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FILING DATE: 2004-11-03
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CL554761 OG ABa000
DU660702 OG ABa001
DW790906 OR ABa011
DW790916 OR ABa019
CW795911 SP Ba011
DW759061 ASWGZ280.
BZ692135 SP Ba001
DW759063 ASWGZ280.
BZ692135 SP Ba001
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BZ692135 SP Ba001
DW868335 BW868335
BW805921 BW805921
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N71028 YZ3011 £1
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AUTHORS
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KEYWORDS
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CZ820350
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                                                                                                                                                                                                                                                                                OMAP (Oryza Map Alignment Project) - Arizona Genomics Institute
Unpublished (2005)
Contact: Rod A. Wing
                                                                                                                                                                                                                                                                                                                                                           Bukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta, Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae, BEP clade, Ehrhartoideae, Oryzeae, Oryza.

1 (bases 1 to 311)
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OC_Ba0198H18.r OC_Ba Oryza coarctata genomic clone
3', genomic survey sequence.
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Seq primer: CAC TCA TTA GGC ACC CCA
Class: BAC ends.
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                                                                                                                                                                 FORWARD: TAA TAC GAC TCA CTA TAG GG
BACKWARD: CAC TCA TTA GGC ACC CCA
                                                                                                                                                                                           Email: rwing@genome.arizona.edu
PCR PRimers
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Rao,K., Luo,M., Jetty,R., Kudrna,D., Muller,C., S
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Fax: 520 621 1259
                                                                                                                                                                                                                                            University of Arizona
Forbes Building Room 303, Tucson,
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          /clone="OC Ba0198H18"
/tissue_type="leaves"
/dev_stage="mature"
/lab_host="DH108"
                                                  /organism="Oryza coarctata"
/mol_type="genomic DNA"
/db_xref="taxon:77588"
/clone="OC_Ba0198H18"
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clone_lib="OC
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CZ178899 MIAA-8A04
AG325093 MUS muscu
CF208340 CAB20003
CF287058 AGENCOURT
AL302994 Tetraodon
CV214664 EST874374
DV006952 CMB243-H0
CC023557 EST819650
AW155158 mg4600026
CW929167 EDCBUSSTR
DR140280 49191388
AW626734 SWODAFCAP
CB832373 SWMDmfCAV
AW651879 SWMDmfCAV
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A1058040 SWAMCAC12
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CB856215 ri49f02.y
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BI783608 kh35e02.y
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Soderlund,C. a
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le OC_Ba0198H18
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RESULT 3
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1 (bases 1 to 470)

Xu, X., Huang, J., Xu, Z., Qian, B., Zhu, Z., Yan, Q., Cai, T., Zhang, X., Xiao, H., Qu, J., Liu, F., Huang, Q., Cheng, Z., Li, N., Du, J., Hu, W., Shen, K., Lu, G., Fu, G., Zhong, M., Xu, S., Gu, W., Huang, W., Zhao, X., Hu, G., Gu, J., Chen, Z. and Han, Z.

Insight into hepatocellular carcinogenesis at transcriptome level by comparing gene expression profiles of hepatocellular carcinoma with those of corresponding noncancerous liver Proc. Natl. Acad. Sci. U.S.A. 98 (26), 15089-15094 (2001)
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CZ802927 579 bp DNA linear GSS 26-JUL-2005 C Ba0175A06.f OC Ba Oryza coarctata genomic clone OC Ba0175A06 5', genomic survey sequence. CZ802927 CZ802927 GI:71242780 GSS.
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Chinese National Human Genome Center at Shanghai
351 Guo Shoujing Road, Zhangjiang Hi-Tech Park, Pudong, Shanghai
201203, P. R. China
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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lone is available at CHGC
Location/Qualifiers
                                                                                                                                                                                                                                                                                    /note="Vector: pBluescript sk(-); Site_1: EcoRI; Site_2:
                                                                                                                                                                                                                                                                                                                             /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="GKBAJH02"
/clone="GKBAJH02"
/tissue_type="hepatocellular carcinoma"
/dev_stage="Adult"
/lab_host="SOLR"
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                                                                                                                                                                                                            Score 18; DB
Pred. No. 12;
4; Mismatches
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Pred. No. 12;
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CZ820348
LOCUS
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AUTHORS
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11 UGAGCCGUUCGCGGCGAA 28
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OC_Ba0198H17.r OC_Ba Oryza
3', genomic survey sequence.
CZ820348
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BACKWARD: CAC TCA TTA GGC ACC CCA
Plate: 0175 row: A column: 06
Seg primer: TAA TAC GAC TCA CTA TAG
Class: BAC ends.
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; BEP
clade; Ehrhartoideae; Oryzeae; Oryza.
University of Arizona
Forbes Building Room 303, Tucsor
Tel: 520 626 9595
Fax: 520 621 1259
Email: rwing@genome.arizona.edu
                                                                                                                                                                                                                                                                                                               Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; BEP clade; Ehrhartoideae; Oryzeae; Oryza.
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Oryza coarctata
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Arizona Genomics Institute
University of Arizona
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1 (Dases 1 to 579)

Kim, H., Collura, K., Wissotski, M., Byrne, M., Stum, D., Smart, D., Kim, H., Collura, K., Wissotski, M., Byrne, M., Stum, D., Smart, D., Kim, H., Collura, C., Soderlund, C. and
                                                                                                                                                                      OMAP (Oryza Map Alignment Project) - Arizona Genomics Institute Unpublished (2005)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CZ820348.1 GI:71260201
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Fax: 520 621 1259
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                                                                                                                         Contact: Rod A. Wing
Arizona Genomics Institute
                                                                                                                                                                                                                                            Rao, K., Luo, M., Jetty, R.,
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m, H., Collura, K., Wissotski, M., Byrne, M.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /dev_stage="mature"
/lab_host="DH10B"
/clone lib="OC_Ba"
/note="Vector: <u>P</u>AGIBAC1; Site_1: HindIII; Site_2: HindIII"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mol_type="genomic DNA"
/db_xref="taxon:77588"
/clone="OC_Ba0175A06"
/tissue_type="leaves"
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__Ba Oryza coarctata genomic clone
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Pred. No. 13;
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                                                                          Tucson,
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Muller,C., Soderlund,C.
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e OC__Ba0198H17
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KEYWORDS
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CZ804464
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Oryza coarctata
Oryza coarctata
Bukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta;
Spermatophyta, Magnollophyta, Lillopsida, Foales, Poaceae, BEP
Clade, Ehrhartoideae, Oryzeae, Oryza.
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CZ80177B01.f OC Ba Oryza
5', genomic survey sequence.
CZ804464
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University of Arizona
Porbes Building Room 303, Tucson, AZ 85721-0036, USA
Tel: 520 626 9595
Pax: 520 621 1259
                                                                                                                                                                                                                                                                Seq primer: TAA
Class: BAC ends.
                                                                                                                                                                                                                                                                                  FORWARD: TAA TAC GAC TCA CTA TAG GG
BACKWARD: CAC TCA TTA GGC ACC CCA
Plate: 0177 row: B column: 01
Seq primer: TAA TAC GAC TCA CTA TAG
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Seq primer: CAC TCA TTA GGC ACC CCA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Rod A. Wing
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                                                                       /mol type="genomic DNA"
/db xref="taxon:77588"
/clone="OC_ Ba0177801"
/tissue_type="leaves"
/dev_stage="mature"
/lab host="pH108"
/clone_lib="OC_Ba"
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/clone_lib="OC_Ba"
/note="Vector: pAGIBAC1; Site_1: HindIII; Site_2: HindIII"
                                                   /clone_lib="OC_Ba"
/note="Vector: pAGIBAC1;
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/clone="OC_Ba0198H17"
/tissue_type="leaves"
                                                                                                                                                                                                           organism="Oryza coarctata"
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 52.9%;

 Mismatches

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 Score 18;
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                                                   Site_1: HindIII; Site_2: HindIII"
13;
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Length 660;
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RESULT 6
CZ878583
LOCUS
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DU808944
LOCUS
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AUTHORS
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Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                         Local
                                                                                                                                                                                                                         437 TGAGCCGTTCGCGGCGAA
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                                                                591 bp DNA linear GSS 13-DEC-2005 OG_ABa0093E05.r OG_ABa Oryza granulata genomic clone OG_ABa0093E05 3', genomic survey sequence.
DU808944
DU808944.1 GI:83604603
Oryza granulata
Oryza granulata
Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
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Seq primer: TAA TAC GAC TCA CTA TAG
Class: BAC ends.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Rod A. Wing
Arizona Genomics Institute
University of Arizona
Forbes Building Room 303, Tucson,
Tel: 520 626 9595
Fax: 520 621 1259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oryza coarctata (Porteresia coarctata)
Oryza coarctata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OC_Ba0280C17.f OC_Ba Oryza coarctata genomic clone OC_Ba0280C17
5', genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished
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Rao,K., Luo,M., Jetty,R., Kudrna,D., Muller,C., Sode)
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BACKWARD: CAC TCA TTA GGC ACC CCA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: rwing@genome.arizona.edu
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                                                                                                                                                                                                                                                                                                                                                                      /dev stage="mature"
/lab_host="DH10B"
/clone_lib="OC_Ba"
/note="Vector: pAGIBAC1; Site_1: HindIII; Site_2: HindIII"
                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mol_type="genomic DNA"
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/clone="OC_Ba0280C17"
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83.3%; Pred. No.
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Matches
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Contact: Scott A. Jackson
Jackson Laboratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15;
                                                                                                   Arizona Genomics Institute
University of Arizona
Forbes Building Room 303, Tucson,
Tel: 520 626 9595
Fax: 520 621 1259
                                                                                                                                                                                                                                                                                                                                                                                   CL554761
CL554761.1 GI:47632656
GSS.
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OB_Ba0004E01.f OB_Ba Oryza brachyantha genomic clone
OB_Ba0004E01 5', genomic survey sequence.
FORWARD: TAA TAC GAC TCA CTA TAG GG
BACKWARD: CAC TCA TTA GGC ACC CCA
Insert Length: 145000 Std Error: (
Plate: 0004 row: E column: 01
                                                                                                                                                                                                       OMAP Project
Unpublished (2004)
                                                                   PCR PRimers
                                                                                   Email: rwing@genome.arizona.edu
                                                                                                                                                                                        Contact: Rod A. Wing
                                                                                                                                                                                                                                           Kim,H., Yu,Y., Stum,D.,
Kudrna,D., Muller,C., H
                                                                                                                                                                                                                                                                                                            Eukaryota; Viridiplantae; Streptophyta; En
Spermatophyta; Magnoliophyta; Liliopsida;
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Seg primer: CAC TCA TTA GGC ACC
Class: BAC ends.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: sjackson@purdue.edu
Basecalling by phred version 0.020425.c. This sequence was derived
from the raw sequence read by clipping with lucy version 1.19s.
Bases 107-697 of the raw sequence (length 1052) were retained after
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Purdue University
915 W. State St., West Lafayette,
Tel: 7654963621
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SanMiguel, P., Weste
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                                                                                                                                                                                                                                                                                            clade; Ehrhartoideae; Oryzeae; Oryza.
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/mol_type="genomic DNA"
/db_xref="taxon:110450"
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lab host="DH10B T1 phage resistant"
clone_lib="OG_ABa"
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Pred. No.
 Std Error: 0.00
lumn: 01
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SanMiguel, P., Westerman, R., Kim, H., Yu, Y., Wissotski, M., Stum, D., Rao, K., Luo, M., Jetty, R., Kudrna, D., Muller, C., Hatfield, J., Soderlund, C., Wing, R. and Jackson, S.A. OMAP (Oryza Map Alignment Project) - Purdue University Unpublished (2004)
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; BEP
clade; Ehrhartoideae; Oryzeae; Oryza.
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Seq primer: CAC TCA TTA
Class: BAC ends.
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915 W. State St.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Scott A. Jackson Jackson Laboratory
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Oryza granulata
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3', genomic survey
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                                                                                                   /Lissue_type="young leaves"
/lab_host="DH10B T1 phage resistant"
/lab_nost="OG_ABa"
/note="Vector: pAGIBAC1; Site_1: Hin
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/db_xref="taxon:110450"
/clone="OG_ABa0063J17"
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/note="Vector: pAGIBAC1; Site_1: HindIII; Site_2: HindIII"
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88.2%;
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ACCESSION
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KEYWORDS
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SOURCE
ORGANISM
                                                                                                                                                                                                                          RESULT 11
DX290906
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Oryza granulata
Oryza granulata
Bukaryota, Viridiplantae, Streptophyta, Embryophyt
Spermatophyta, Magnoliophyta, Liliopsida, Poales,
Clade, Bhrhartoideae, Oryzeae, Oryza.
Oryza ridleyi
Oryza ridleyi
Oryza ridleyi
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; BEP
clade; Ehrhartoideae; Oryzeae; Oryza.
1 (bases 1 to 795)
                                                                                                                                                                      DX290906
OR ABa0194J18.f OR ABa Oryza
5', genomic survey sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bmail: sjackson@purdue.edu
HindIII (presumptive cloning site) found. Basecalling by phred
HindIII (presumptive cloning site) found. Basecalling by phred
version 0.020425.c. This sequence was derived from the raw sequenc
read by clipping with lucy version 1.19s. Bases 31-793 of the raw
sequence (length 1065) were retained after clipping.
Plate: 0110 row: K column: 10
Seq primer: TAA TAC GAC TCA CTA TAG GG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GSS.
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915 W. State St., West Lafayette,
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DX010184
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /tissue_type="young leaves"
/lab_host="DH10B T1 phage resistant"
/clone_lib="OG_ABa"
/note="Vector: pAGIBAC1; Site_1: HindIII; Site_2: HindIII"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   organism="Oryza granulata"
/mol_type="genomic DNA"
/db_xref="taxon:110450"
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Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD
clade; Panicoideae; Andropogoneae; Sorghum.
1 (bases 1 to 904)
2 Andropogoneae; Andropogoneae; Sorghum.
                                                                                                                                                                                                                                                                                           Bowers, J.E., Arias, M.A., Asher, R., Avise, J.A., Ball, R.T., Brewer, G.A., Buss, R.W., Chen, A.H., Edwards, T.M., Betill, J.C., Exum, H.E., Goff, V.H., Herrick, K.L., Steele, C.L., Karunakaran, S., Lafayette, G.K., Lemke, C., Marler, B.S., Masters, S.L., McMillan, J.M., Nelson, L.K., Newsome, G.A., Nwakamma, C.C., Odeh, R.N., Phelps, C.A., Rarick, E.A., Rogers, C.J., Ryan, S.P., Slaughter, K.A., Soderlund, C.A., Tang, H., Wing, R.A. and Paterson, A.H. Comparative physical mapping links conservation of microsynteny to chromosome structure and recombination in grasses proc. Natl. Acad. Sci. U.S.A. 102 (37), 13206-13211 (2005)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CW785911 904 bp DNA linear GSS SP_Ba0016N09.r SP_Ba Sorghum propinquum genomic clone SP_Ba0016N09 3', genomic survey sequence.
             FORWARD: TAA TAC GAC TCA CTA TAG GG
BACKWARD: CAC TCA TTA GGC ACC CCA
Plate: 0016 row: N column: 09
Seg primer: CAC TCA TTA GGC ACC CCA
                                                                                                                                               University of Arizona
Forbes Building Room:
Tel: 520 626 9595
Fax: 520 621 1259
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University of Arizona
Forbes Building Room 303, Tucson,
Tel: 520 626 9595
Fax: 520 621 1259
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Plate: 0194 row: J column: 18
Class: BAC ends.
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Rao,K., Luo,M., Jetty,R., Kudrna,D., Muller,C., Soderlund,C.
Class: BAC ends
                                                                                                  PCR PRimers
                                                                                                                      Email: rwing@genome.arizona.edu
                                                                                                                                                                                                                                                        Contact: Rod A. Wing
                                                                                                                                                                                                                                    Arizona Genomics Institute
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/mol_type="genomic DNA"
/db_xref="taxon:83308"
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/note="Vector: pAGIBAC1; Site_1: HindIII; Site_2: HindIII"
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/lab_host="DH10B"
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                                                                                                                                                                                                                                                                                                                                                                                                                     Email: PMRichardson@lbl.gov; delong@mit.edu
North Pacific Subtropical Gyre (Hawaii) picoplankton genomic fosmid
DNA library prepared from marine picoplankton in the less than 1.6
um, greater than 0.2 um fraction. Sample Date: 10/6/2002
Coordinates: 22.45 N, 158 W Depth 200 m Temperature: 18.53 C
Salinity: 35.04 psu Oxygen: 198.8 umol/kg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US DOE Joint Genome Institute
US DOE Joint Genome Institute
2800 Mitchell Drive B100, Walnut Creek, CA 94598-1698,
Tel: 617-253-5271
Fax: 617-253-2679
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Science (2006) In press
Contact: Susan Lucas, Alex Copeland, Sam Pitluck,
Kerrie Barry, Tijana Glavinadelrio, David Bruce, I
and Edward DeLong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Comparative genomics reveals ecological trends in stratified microbial communities in the ocean's interior
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 955)
DeLong, B.F., Preston, C.M., Mincer, T., Rich, V., Hallam, S.J.,
Frigaard, N.U., Martinez, A., Sullivan, M., Edwards, R., Chisholm, S.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and Karl, D.M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     uncultured marine microorganism HF200_10-06-02 uncultured marine microorganism HF200_10-06-02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 unclassified sequences; environmental samples.
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                                                                                                                                                                                                                                                                                                                                                                                                        fosmid ends.
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/clone lib="HF200 10-06-02"
/clone lib="HF200 10-06-02"
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/note="Vector: pCCIFOS; North Pacific Subtropical Gyre
/note="Vector: pCCIFOS; North Pacific Subtropical Gyre
/framily picoplankton genomic fosmid DNA library prepared
from marine picoplankton in the less than 1.6 um, greater
than 0.22 um fraction. Picoplankton collected at 200 m
depth on 10/6/2002, Coordinates: 22.45 N, 158 W Depth 200 m
depth on 10/6/2002 Coordinates: 22.45 N, 158 W Depth 200 m
Temperature: 18.53 C Salinity: 35.04 psu Oxygen: 198.8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone="SP_Ba0016N09"
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/clone_lib="SP_Ba"
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HindIII; Paterson lab BAC library (HindIII)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /db_xref="taxon:132711"
/clone="SP_Ba0016N09"
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/mol_type="genomic DNA"
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                                                                                                                                                                                                                                                                                                                                     organism="uncultured marine microorganism HF200_10-06-02"
                                                                                                                                                                                                                                                                                                                                                                               ocation/Qualifiers
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76.5%;
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Paul Richardson
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KEYWORDS
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BW868335
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273 GATGAGCCGTTCGCGGC 289
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                                                                                                             BW868335 Amphioxus Branchiostoma floridae unpublished cDNA library, neurula whole animal Branchiostoma floridae cDNA clone bfne085101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Arizona Genomics Institute
University of Arizona
Biological Sciences West, 448A, P.O. Box 210088, Tucson, 85721-0088, USA
Tel: 520 626 3967
Fax: 520 621 9288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequencing of Sorghum BAC ends.
http://genome.arizona.edu/stc/sorghum
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sorghum propinquum
Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD
clade; Panicoideae; Andropogoneae; Sorghum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SP Ba0016N09.r SP Ba Sorghum propinquum genomic clone SP Ba0016N09 3', genomic survey sequence.

BZ692135
Branchiostoma floridae (Florida lancelet) Branchiostoma floridae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Plate: 0016 row: N column: Seq primer: gta aaa cga cgg Class: BAC ends.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 (bases 1 to 1299)
Wing,R., Yu,Y., Kim,H.R., Collura,K., Pries,G., Currie,J.,
Soderlund,C. and Hatfield,J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BZ692135.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: rwing@genome.arizona.edu
PCR PRimers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Rod Wing
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                                                         BW868335.1
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                                                                                               mRNA sequence.
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COMMENT
Search completed: August 28, 2006, 10:29:08 Job time : 2336 secs
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                                                                                                 45 TCGCGGCGAAACGAAG 60
                                                                                                                                                 19 UCGCGGCGAAACGAAG 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 359)
Yu,J., Holland,L.Z., Shin-i,T., Kohara,Y., Satou,Y. and Satoh,N. Expressed genes in Branchiostoma floridae
Unpublished (2005)
Contact: Tadasu Shin-i
Center For Genetic Resource Information
National Institute of Genetics
1111 Yata, Mishima, Shizuoka 411-8540, Japan
Tel: 81-559-81-6855
Fax: 81-559-61-6855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: tshini@genes.nig.ac.jp
If you want to have a cDNA clone for this EST or if you have any
questions, please send an e-mail to Nori Satoh
(satch@ascidian.zool.kyoto-u.ac.jp) and its cc to Yutaka Satou
(yutaka@ascidian.zool.kyoto-u.ac.jp).
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bukaryota; Metazoa; Chordata; Cephalochordata; Branchiostomidae; Branchiostoma.
                                                                                                                                                                                                                                                                                                                /organism="Branchiostoma floridae"
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/clone="bfne085101"
/tissue_type="whole animal"
/dev_stage="neurula"
/clone_lib="Amphioxus Branchiostoma floridae unpublished
cDNA library, neurula whole animal"
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Fauchet,C.R.J.
Fixing unit with an end imprint in a threaded terminal portion
Patent: US 6632057-A 100 14-OCT-2003;
GPI Aerospace; Paris;
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/mol_type="unassigned RNA"
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Fauchet,C.R.J.
Fixing unit with an end i
Patent: US 6632057-A 98 1
GFI Aerospace; Paris;
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Fauchet C.R.J.
Fixing unit with an end imprint in a Pixing unit with an end imprint in a Patent: US 6632057-A 92 14-OCT-2003; GFI Aerospace; Paris;
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Sequence 92 from patent US
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Pred. No. 7.7e-06;
4; Mismatches 0;
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Fauchet, C.R.J.
Fixing unit with an end in
Patent: US 6632057-A 99 1-
GFI Aerospace; Paris;
FRX;
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Patent: JP 2002542805-A 57 1:
Patent: JP 2002542805-A 57 1:
UNIVERSITY OF FLORIDA
OS Artificial Sequence
PN JP 2002542805-A/57
PD 17-DEC-2002
PF 28-APR-2000 JP 200061540
PR 30-APR-1999 US 60/131
PI ALFRED S LEWIN, NICHOLAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PF 28-APR-2000 JP 2000615402
PR 30-APR-1999 US 60/131942
PI ALFRED S LEWIN, NICHOLAS MUZYCZKA, WILLIAM W HAUSWIRTH , CHRISTIAN TESCHENDORF,
PI CORTINA BURGER
PC C12N16/00 1000
                                                                                                                                Fixing unit with an end
Patent: US 6632057-A 90
GFI Aerospace; Parin:
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AR407997
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Description of Artificial Sequence: SYNTHETIC PEPTIDE
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1 (bases 1 to 34)
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JP 2002542805-A/57.
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Adeno-associated virus-delivered
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                                                                                                                                                                                        Unclassified.
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ACUGAUGAGCCGUUCGCGGCGAAAC 30
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/mol_type="genomic RNA"
/db_xref="taxon:32630"
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                                                                           /organism="unknown"
/mol_type="unassion«
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Pred. No. 0.00015;
4; Mismatches 0
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Fauchet, C.R.J.
Fixing unit with an end
Patent: US 6632057-A 95
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Fauchet, C.R.J.
Frauchet, T.R.J.
Patent: US 6632057-A 91 14-OCT-2003;
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/mol_type="unassigned
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Fauchet, C.R. J.
Fixing unit with an end i
Patent: US 6632057-A 96 1
GFI Aerospace; Paris;
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Sequence
AR408004
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of use
Patent: WO 0066780-A 57 09-NOV-2000;
University of Florida (US)
Location/Qualifiers
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21; Conserv
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Fauchet, C.R.J.
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                                                                                 Muzyczka,N.,
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Pred. No. 0.00
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                                                                                    Teschendorf, C.
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UNIVERSITY OF FLORIDA
OS Artificial Sequence
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Description of Artificial Sequence: SYNTHETIC OLIGONUC
Key Location/Qualifiers
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                      synthetic construct synthetic construct
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ALFRED S LEWIN,NICHOLAS MUZYCZKA,WILLIAM W HAUSWIRTH
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            sequences; artificial sequences.
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Location/Qualifiers
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/db_xref="taxon:32630"
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